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    GenCore version 5.1.6 (c) 1993 - 2006 Compugen Ltd.
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  Adi80376
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 Human car
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Bacillus
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HUMAN T C
Human TRF
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AAY90110
AAY87685
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Aar31678768

Aar951942

Aab2894110

Aay90110

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1 HIV 9p120
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7 HLA bindi
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                                                                                                                                                                                                                                                                               CC -map) composition. The HMW-map composition was isolated from a D. farinae CC homogenate by gel filtration, with each fraction being analysed for the presence of proteins that bound to IgE present in mite-allergic dog CC antisera. The HMW-map composition comprises mapA (a 109 kD protein) and CC mapB (98 kD). Mite allergenic proteins and peptides, and nucleic acids cencoding them, may be used in therapeutic compositions to modify an CC enimal's hypersensitivity reaction to mite allergens. Animals that may be treated include mammals and birds, especially feilnes, canines, equines, commans, other pets, and work or domestic animals. The proteins or CC fragments may also be used to diagnose allerges via a skin test. The proteins and peptides can also be used to raise antibodies, which have a corriety of potential uses. For example, they can be used as vaccines to passively immunise animals against dust mite hypersensitivity, as CC passively munise animals against dust mite hypersensitivity, as comparing the proteins of potential uses. For example, they can be used as vaccines to passively immunise animals against dust mite hypersensitivity, as constitute controls in test kits and as tools to recover desired dust mite allergens from a mixture of proteins
                                                                                                                      Matches
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Best Local :
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86
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mite allergen protein; map; high molecular weight; HMW-map; allergy; house dust mite; IgE; immunoglobulin E; allergen; mapA; mapB; hypersensitivity reaction; therapy; treatment; diagnosis; human; fel
                                                                                                                                                                                                                                         Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Page 69; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel high molecular weight Dermatophagoides nucleic acid polypeptides used to modify an animals' hypersensitivy to mite allergens.
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13-MAY-1998;
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llarity 100.0%;
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98US-0098909P.
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Pred. No. 0.00013;
; Mismatches 0;
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AAU96320
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ID AAU96320
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XX Der
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XX Der
XX W mite
XX W mite
XX W 14-1
PD 21-1
PF N MCC
XX HPI
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disease. Antibodies that bind to Der HMW-map are useful for inhibiting binding of proteins to IgE, to prevent immunocomplex formation, thus reducing hypersensitivity responses to mite allergens, and as vaccines against mite allergen hypersensitivity. Sequences AAU96314-AAU96342 represent Der HMW-map polypeptides of the invention
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Human cartilage glycoprotein 39 derived peptide #21

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a peptide which resembles or mimics an epitope present on human cartilage glycoprotein 39 (HC gp-39), an autoantigen in rheumatoid arthritis. The invention relates to peptides consisting of 16-55 amino acid residues comprising at least one of the following 19 sequences: LVCYYTSWS, FLCTHIYS; INTSTANTS; LKTLLSYGG; FRGOEDLAML; LYPGRADKG; YLAXISCH; LLDETSIMTY; FISIMTYDF; FRGOEDLAML; LYPGRADKG; LAYYEICDF; LRGATVHTT; YLKURQLAG; LAGAMVWAL; WALDLDDF; or LDLDDFQGS. They can be used medically in antigen specific immunosuppressive therapy, particularly the treatment of T-cell mediated destruction of articular cartilage in autoimmune diseases (e.g. rheumatoid arthritis). They can also be used to detect activated autoreactive T cells in an individual. The peptides have a specific confect on the autoreactive T cells, thus leaving the other components of the immune system intact, unlike the non-specific suppressive effect of the immune system intact, unlike the non-specific suppressive effect of the immune system intact, unlike the non-specific suppressive effect of the monauppressive drugs, and do not cause toxic side effects. The peptides are predominantly recognised by autoreactive T cells from relating the other components.
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Best Local
Major Histocompatibility Complex; MHC; CDB; CD4; immunosuppressive; antidiabetic; antirheumatic; antiarthritic; neuroprotective; antiinflammatory; gene therapy; T cell response; autoimmune disorder; graft-versus-host disease; graft rejection; rheumatoid arthritis; celiac disease; dermatitis herpetiformis; diabetes; multiple sclerosi. Crohn's disease; Psoriasis; human leukocyte antigen.
                                                                                                                                                                                                                               Human leukocyte antigen A sub-type peptide region 110-130 group
                                                                                                                                                                                                                                                                                                       22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                             ADI80376 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                       (first entry)
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Pred. No. 29;
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                                     multiple sclerosis;
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RESULT 5
AAY99022
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AC AAYS
AC AAYS
DT 07-1
XX
DE HLA
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AAY99022 standard; peptide;

13

07-AUG-2000

(first entry)

HLA class II binding antigen epitope peptide #211

밁 S

10

DGKNYIALNEDL DKONYLALVREL

21

Matches Query Match Best Local :

Similarity 6; Conserv

Conservative

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Score 31; DB 5; Pred. No. 2.1e+02; 3; Mismatches 3

Length 21; Indels

0

Gaps

0

48.4**%**; 50.0**%**;

Sequence 21

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CC Complex (MHC) molecule or a nucleic acid encoding the modified MHC CC molecule. The binding of the MHC molecule of a selected type to CDB or CC CD4 is inhibited, but it can present the same peptide or peptides as CC unmodified molecules of the MHC type, which is for use in medicine. The CC invention further comprises: inhibiting the activity of T cells against a CC cell presenting molecules of the MHC type, which is for use in medicine. The CC present modified molecules of the MHC type, by causing the cell to CC present modified molecules of the MHC type; a cell, which presents CC molecules of a selected MHC type, by causing the cell to CC can present the same peptide or peptides as unmodified molecules of the CC can present the same peptide or peptides as unmodified MHC molecules or its complementary sequence; a vector comprising the molecule acid molecule encoding the modified MHC molecule has the CC following activities: immunosuppressive, antidiabetic, antirheumatic, antiarthritic, neuroprotective, and antiinflammatory. The nucleic acid cencoding the modified MHC molecule, as well as the nucleic acid condoing the modified MHC molecule, as well as the nucleic acid cencoding the modified MHC molecule, as well as the nucleic acid condoing the molecule, are useful in the manufacture of a medicament for companion the modified MHC molecule, as well as the nucleic acid condogenous or exogenous etiology), graft-versus-host disease or graft crejection. These disorders also include rheumatoid arthritis, celiac condimentation herpetiformis, diabetes, multiple sclerosis, Crohn's colypeptide relating to the modified MHC molecule of the invention.
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New modified Major Histocompatibility Complex (MHC) molecule or nucleic acid encoding the MHC molecule, useful in medicine, particularly useful for treating autoimmune disorders, e.g. graft-versus-host disease or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-723521/78
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RESULT 6
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                                                                                                                                                                                                                                                                                                                             CC comprising a unit dose form of a peptide, or analogue, compressing an CC epitope selected from those represented by peptides ANY98812-Y99339 which are derived from various antigens for various human leucocyte antigen CC class DR molecules, representative of the world wide population. The peptide/analogue binds to an HLA class II molecule at an IC-50 of less CC than or equal to 1,000 nM. The pharmaceutical focuses the immune response CC than or equal to 2,000 nM. The pharmaceutical focuses the immune response CC chronic vital diseases and cancer. Examples of diseases that can be CC treated using the peptide containing pharmaceutical include autoimmune CC diseases (rheumatoid arthritis, multiple sclerosis, and myasthenia CC gravis), allograft rejection, allergies, lyme disease, hepatitis, post-CC streptococcal endocarditis or glomerulonephritis and food CC hypersensitivities. The peptide epitopes can be used to enhance immune CC diseases which can be treated using immunogenic mixtures include prostate CC cancer, hepatitis B, hepatitis C, AIDS, renal carcinoma, cervical CC carcinoma, lymphoma, and condyloma acuminatum. The peptides may also be used to make monoclonal antibodies useful as potential diagnostic or therapeutic agents. The peptides may also be useful as diagnostic or therapeutic agents. The peptides may also be useful as diagnostic corresponse which as substantial risk of developing chronic infection. CC me selection of appropriate T and B cell epitopes should allow the development of pathogens which are characterized by high sequence correlations of pathogens which are characterized by high sequence
                                                                                                                                                                                                                          Query Match
Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  glomerulonephritis;
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                 ADW33519;
                                                      ADW33519 standard; peptide;
                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a new pharmaceutical composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New compositions containing immunogenic peptide epitopes for various class II DR molecules useful for inducing helper {\tt T} cell response.
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85.7%;
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Pred. No. 1.9e
0; Mismatches
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1.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                           composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an AZ restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnosing viral diseases and cancer. The peptide epitopes are useful as diagnostic agents for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA;
Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T cell; MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA; viral disease; cancer.
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                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 52-379; 382pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            binding motifs for HLA, us viral diseases and cancer.
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20-APR-2001; 2001US-0285624P
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                                                                HLA binding
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                                                                                                                                                                ADW34752 standard; peptide; 13
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                                                               epitope #5502
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85.7%;
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Pred. No. 1.9e+02;
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Best Local
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Dac gI;
                                 29-SEP-1994
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                                                                                        WO9421675-A2
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01-JUN-1995
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viral diseases and cancer.
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20-APR-2001;
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2001US-0285624P.
                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                      protein allergen; Lol pI; ryegrass pollen allergen;
; Phl pI.
                                                                                                                                                                                                                                                                                        protein allergen internal sequence LPI-16.10
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85.7%;
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Pred. No. 1.9e+02;
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Best Local
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                                            New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing viral diseases and cancer.
                                                                                                                                                                                                                                                                                                     Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA; viral disease; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated peptide(s) of Lol or p I, major protein allergen of species Lolium perenne. - useful for diagnosis and treatment of sensitivity to rye-grass pollen allergen.
                     Claim 1; Page 52-379; 382pp; English.
                                                                                            WPI; 2003-441519/41.
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20-APR-2001;
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                                                                                                                                                                                                                                                                               Unidentified
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5; Conserv
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2001US-0285624P
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Pred. No. 2.2e
3; Mismatches
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The invention relates to a composition comprising

at least one

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CC from 30 lists given in the specification. Also disclosed is a method for confidence in a cytotoxic T cell response against a pre-selected antigen in a gretient expressing a specific MFC class I allele by contacting cytotoxic T cell response against a pre-selected antigen in a CC patient expressing a specific MFC class I allele by contacting cytotoxic CC composition comprises an epitope that is joined by an amino acid linker. CC chapter is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an A2 CC restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically CC claimed are peptides having allele-specific binding motifs for HLA. The CC diagnositions and methods are useful for preventing, treating or CC diagnosing viral diseases and cancer. The peptide epitopes are useful as CC diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-CC ADW37745 represent epitopes of the invention as given in Tables 2-31.
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having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is
                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                         binding motifs for HLA, us
viral diseases and cancer.
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ling motifs for HLA, useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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DKEGFLNKVRE

12

Matches Query Match Best Local

6 Similarity

Conservative

46.9%;

Score 30; I Pred. No. 3

DB .1e+02;

2

Length

20

Mismatches

Indels

0

Gaps

0

DKONYLALVRE 11

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RESULT 11
AAW97099
ID AAW977
XX AAW97
XX AAW97
XX DT 30-AE
XX Therm
KW Therm
KW In vi
XX Pyroc
XX AAW9:
CC AAW9:
CC AAW9:
CC AAW9:
CC AGtc:
XX Seque
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnosing viral diseases and cancer. The peptide epitopes are useful as diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
                              AAW97099-104 represent peptides isolated from the cell fluid obtained after rupture of Pyrococcus furiosus DSM3638. The peptides are derived from thermostable polypeptide factors which bind to, and promote the DN synthesis activity of DNA polymerase. The polymerase related factors cabe used to provide more efficient in vitro DNA synthesis and amplification systems (e.g. for polymerase chain reaction) by using the factors in conjunction with a DNA polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW97099;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW97099 standard; peptide; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bound to
 Sequence
                                                                                                                                                                   Example
                                                                                                                                                                                                  Thermostable polypeptide for improvement of DNA
                                                                                                                                                                                                                                                   WPI; 1999-095751/08
                                                                                                                                                                                                                                                                                    Uemori
                                                                                                                                                                                                                                                                                                                                                   21-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                26-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                  24-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pyrococcus furiosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide from cell fluid obtained after rupture of P. furiosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                  (TAKI ) TAKARA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in vitro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                    T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 KONYLALVRE 11
                                                                                                                                                                   3; Page 34; 177pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KONTLAFVKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16
     80
                                                                                                                                                                                                                                                                                  Sato Y,
     ΑA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLA molecule on the antigen-presenting cell, where when an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptide factor; DNA synthesis synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                    SHUZO CO LTD.
                                                                                                                                                                                                                                                                                                                                                   97JP-00320692
                                                                                                                                                                                                                                                                                                                                                                97JP-00187496
                                                                                                                                                                                                                                                                                                                                                                                                  98WO-JP002845
                                                                                                                                                                                                                                                                                  Fujita T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46.9%;
                                                                                                                                                                                                Score 30; DB
Pred. No. 2.4e
2; Mismatches
                                                                                                                                                                                                                                                                                  Miyake K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          . 2.4e+02;
2;
                                                                                                                                                                                                                                                                                  Mukai H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 activity; DNA polymerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                    Asada
                                                                                                                                                                                                                                                                                    ζ,
                                                   by using the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                    Kato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                   can
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RESULT 13
AAW45121
ID AAW45
XX
AC AAW45
AC AAW45
XX
DT 28-AF
XX
DE Human
                                                                                                                                                                                                                    Query Match
Best Local Similarity
''--'hes 5; Conservi
                                                                                                                                                                         밁
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                                                                                                                                                                                                                                                                                                  RESULT 12
AAM42730
                                                                                                                                                                                                                                                                                                                       protein or receptor and are thus useful as reagents and drugs, and as lead ligands to facilitate drug design and development. They are useful as tools for functional genomic studies, reagents for the configuration of high-throughput screens, as a starting point for medicinal chemistry manipulation, for peptide mimetics and as therapeutic agents. The analysis and acquisition of peptide sequences facilitates understanding of protein-protein interactions. The method allows for analysis of an entire database at a time, thus overcoming sampling problems. The set of complementary peptides includes both intermolecular (between proteins) and intermolecular (within a protein) sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is one of a large number of complementary peptide ligands generated from Mycoplasma genitalium genome sequences. These specific complementary peptides interact with their relevant target proteins encoded by the microbial genome. They are capable of antagonising or agonising specific interaction of a protein with another
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Complementary peptide ligands as reagents and drugs for drug discovery programs and as lead ligands to facilitate drug design and development, are generated from microbial genome sequences.
Human cartilage glycoprotein 39 derived peptide #22
                                28-APR-1998
                                                                                           AAW45121 standard; peptide; 16
                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-514238/56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Roberts GW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mycoplasma genitalium; complementary peptide; ligand; protein-protein interaction; drug design; intermolecular; intramolecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAM42730 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 55; 161pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mycoplasma genitalium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mycoplasma genitalium intermolecular complementary peptide, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-OCT-2001
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                                                                                                                                                                                                     5 YLALVRELK 13
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                                                                                                                                                                        FLSLVKEVK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Heal JR;
                                                                                                                                                                                                                                    Conservative
                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99GB-00029466
                                                                                                                                                                         9
                                                                                                                                                                                                                                                 45.3%;
55.6%;
                                                                                                                                                                                                                               Score 29; DB 4; Len;
; Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10
                                                                                                                                                                                                                                                                 Length 10
                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                    0,
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Human 18-SEP-2001 AAG95826;

complementary peptide,

SEQ ID

NO: 2020.

(first entry)

Human; complementary peptide; ligand; drug discovery; drug design.

WO200142277-A2

AAG95826 RESULT 14

AAG95826 standard; peptide;

10

맑 S

ONYLALVRELK 13 QHFTTLIKEMK

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The present sequence represents a peptide which resembles or mimics an cepitope present on human cartilage glycoprotein 39 (HC gp-39), an autoantigen in rheumatoid arthritis. The invention relates to peptides consisting of 16-55 amino acid residues comprising at least one of the following 19 sequences: LVCYYTSNS; FLCTHIYS; HISTANIS; LKTLLSVGG; FRCQEDASH; YAVGYMLRL; MLRLGAPAS; LAYYEICDF; LDFISIMTY; FISIMTYDF; CC FRCQEDASH; YAVGYMLRL; MLRLGAPAS; LAYYEICDF; LRGATVHRT; YLKDRQLAG; LAGAMWWAL; WALDLDDF; or LOLDDFQGS. They can be used medically in antigen specific immunosuppressive therapy, particularly the treatment of T-cell mediated destruction of articular cartilage in autoimmune diseases (e.g. rheumatoid arthritis). They can also be used to detect activated autoreactive T cells in an individual. The peptides have a specific immunosuppressive drugs, and do not cause toxic side effects. The peptides are predominantly recognised by autoreactive T cells from rheumatoid arthritis patients, but rarely by those from healthy donors
         Matches
                                       Query Match
Best Local
                                                                                                                               Sequence 16 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide suitable for use in antigen specific immunosuppressive therapy resembles or mimics epitope present on HC gp-39, so inducing systemic immunological tolerance to rheumatoid arthritis auto:antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Boots AMH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Articular cartilage; immunosuppressive therapy; antigen; autoantigen; immunological tolerance; T-cell; human cartilage glycoprotein 39; HC gp-39; rheumatoid arthritis; epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ALKU ) AKZO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9740068-A1
                                   Similarity
         4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Verheijden GFM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 16;
         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOBEL NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96EP-00201106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97WO-EP002051
                                   45.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   82pp; English.
Score 29; DB 2; Le
Pred. No. 3.6e+02;
Pred. No. 3.6e+02;
                                                                 Length 16;
         Indels
      0,
      Gaps
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RESULT 15
ABP68019
ID ABP68
XX ABP68
XX ABP68
XX BBC1
XX BBC1
XX BBC1
XX BBC1
XX BBC1
XX BBC2

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A set of peptide ligands consisting of specific complementary peptides to proteins encoded by genes of the human genome, useful in an assay for screening and identifying of one or more novel peptides which are drug candidates or pro-drugs.
                                                            Example 2; Page 21; 134pp; French.
                                                                                                                                                                                                                                                               Freyssinet G,
                                                                                                                                                                                                                                                                                                                                                                                                                                 19-MAR-2001; 2001FR-00003691.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FR2822157-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacillus thuringiensis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pepsin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacillus thuringiensis; insecticide; toxin; Cry; pepsin cleavage site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacillus thuringiensis toxin Cry related peptide #12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABP68019;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABP68019 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; Page 332; 646pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-408419/43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roberts GW, Heal JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PROT-) PROTEOM LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JUN-2001.
                                                                                                                                                                                                                                                                                                                      (AVET ) AVENTIS CROPSCIENCE SA.
                                                                                                                                                                                                                                                                                                                                                                          19-MAR-2001; 2001FR-00003691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the specification
                                                                                                                 modified Cry protein, useful as insecticide, comprises at least one itional pepsin cleavage site to reduce persistence in mammalian gut.
                                                                                                                                                                                                        2003-002439/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DEERYLAL 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DKQNYLAL 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                         Rang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99GB-00029464.
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                                                                                                                                                                                                                                                            Frutos R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Score 28; DB;
; Pred. No. 3.2e.
2; Mismatches
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3.2e+02;
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present invention describes a modified Cry protein (I) that is

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sensitive to pepsin and comprises at least one additional pepsin cleavage site (PCS). Also described: (a) increasing pepsin sensitivity of Cry proteins by incorporating at least one extra PCS; (b) polynucleotides (II) that encode (I); (c) chimeric genes (CG) that contain a promoter, (III) that encode (I); (d) expression or transformation vector (III) that contains CG; (e) host organism (IV) transformed with (III), also, where the organism is a plant, its parts and seeds; (f) production of (I) by growing (IV); and (g) mono- or polyclonal antibodies (Ab) directed graphist (I). (I) has insecticide activity. (I) can be used as insecticides, particularly where expressed in transgenic plants. (I) are sensitive to enzymes in the digestive tract of mammals, so do not persist in the tract (lack of persistence is required by regulatory authorities for use, in foods, of seeds containing Cry proteins). Extra PCS do not insecticidal activity. (ABV93909 and ABP67997 to ABP68308
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The present invention describes a modified Cry protein (I) that is sensitive to pepsin and comprises at least one additional pepsin cleavage site (PCS). Also described: (a) increasing pepsin sensitivity of Cry proteins by incorporating at least one extra PCS; (b) polynucleotides (II) that encode (I); (c) chimeric genes (CG) that contain a promoter, (II) and terminator; (d) expression or transformation vector (III) that contains CG; (e) host organism (IV) transformed with (III), also, where the organism is a plant, its parts and seeds; (f) production of (I) by growing (IV); and (g) mono- or polyclonal antibodies (Ab) directed against (I). (I) has insecticide activity. (I) can be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABP68026 standard; peptide; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacillus thuringiensis Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacillus thuringiensis; insecticide; toxin; Cry; pepsin cleavage site; pepsin; PCS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacillus thuringiensis toxin Cry related peptide
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                                                                                                                                                                                                      Example 2; Page 21; 134pp; French.
                                                                                                                                                                                                                                          New modified Cry protein, useful as insecticide, comprises at least one additional pepsin cleavage site to reduce persistence in mammalian gut.
                                                                                                                                                                                                                                                                                                   WPI; 2003-002439/01.
                                                                                                                                                                                                                                                                                                                                      Freyssinet G,
                                                                                                                                                                                                                                                                                                                                                                                                            19-MAR-2001; 2001FR-00003691
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                          (AVET ) AVENTIS CROPSCIENCE SA.
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4e+02;
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RESULT 17
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19-JUL-2002;
19-JUL-2002;
03-MAR-2003;
contain a nucleic acid encoding a mutant HSV protein or a nucleic acid encoding the polypeptide sufficient for virus-like particle assembly but devoid of late-domain motifs The nucleic acid is within an HSV genome. The genome is devoid of late domain motifs capable of effecting viral budding. The composition also comprises a compound capable of interfering with the protein-protein interaction between a host cell protein capable of binding a late domain motif and a HSV protein containing a late domain motif is and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV protein containing a late domain motif and a HSV pr
                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to cells displaying herpes simplex virus (HSV) altered budding phenotype which are useful for the manufacture of a medicament for treating HSV infection. The medicament further comprises an adjuvant capable of stimulating cellular immune response e.g. II-2, IL
                                                                                                                                                                                                                                                                                                                                          an adjuvant capable of stimulating cellular immune response e.g. IL-4, IL-12, IL-18 or gamma-interferon. The cells are human cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 14;
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; 2002US-0397477P.
; 2002US-0397479P.
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Pred. No. 4.3e
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
The genome is devoid of late-domain motifs. The nucleic acid is within an HSV genome. The genome is devoid of late domain motifs capable of effecting viral budding. The composition also comprises a compound capable of interfering with the protein protein interaction between a host cell protein capable of binding a late domain motif and a HSV protein containing a late domain motif. The composition further comprises another HSV protein or its immunogenic fragment, and/or a nucleic acid encoding the other HSV protein or the immunogenic fragment. The present sequence is a mutant HSV peptide, derived from the wild-type pentide and antigory.
                                                                                                                                                                                                                                                                                                                                                                                                                         19-JUL-2002;
19-JUL-2002;
19-JUL-2002;
                                                                                                                               The present invention relates to cells displaying herpes simplex virus (HSV) altered budding phenotype which are useful for the manufacture of medicament for treating HSV infection. The medicament further comprises an adjuvant capable of stimulating cellular immune response e.g. IL-2, I-4, IL-18 or gamma-interferon. The cells are human cells and contain a nucleic acid encoding a mutant HSV protein or a nucleic acid encoding the polypeptide sufficient for virus-like particle assembly but
                                                                                                                                                                                                                                                                             Use of cells displaying herpes simplex virus (HSV) altered budding phenotype for the manufacture of a medicament for treating HSV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunogenic fragment, and/or a nucleic acid encoding the other protein or the immunogenic fragment. The present sequence is a peptide, derived from the wild-type peptide ADJ38601, used to the invention.
                                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                          WPI; 2004-123282/12
                                                                                                                                                                                                                                                                                                                                                     Morham
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                                                                                                                                                                                                                                                                                                                                                                                  MYRIAD GENETICS INC
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                                                                                                                                                                                                                                                                                                                                                     Zavitz
                                                                                                                                                                                                                                                                                                                                                                                                           ; 2002US-0397265P.
; 2002US-0397477P.
; 2002US-0397479P.
; 2003US-0451903P.
                                                                                                                                                                                                                                                  Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers 7..8
                                                                                                                                                                                                                                                    68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Wild-type residues replaced with
                                                                                                                                                                                                                                                                                                                                                    ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13
                                                                                                                                                                                                                                               74pp;
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                                                                                                                                                                                                                                                                                                                                                     Hobden
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antiviral; late domain motif; mutein; mutant.
                                                                                                                                                                                                                                                  English.
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Pred.
1; Mis
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5.1e+02;
3;
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illustrate
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                                          Query Match
Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                        Gefter ML, I
                                                                                                                                                                                                                                                                                                                                                                  25-MAR-1992;
15-MAY-1992;
15-JAN-1993;
                                                                                                           The sequences given in AAR41975-82 are peptides derived from a human T cell reactive feline protein. These peptides are used in a therapeutic composition which is useful in treating diseases which involve an immur response to a protein antigen. This composition may be used to induce tolerance in a mammal to Dermatophagoides, Felis, Ambrosia, Lolium, Cryptomeria, Alternaria, Alder, Betula, Ouercus, Olea, Artemesia, Plantago, Parietaria, Canis, Blattella, Apis, Periplaneta and to autoantigens in humans. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                       New peptide(s) for inducing of an allergen administered cats, bees, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; T cell; reactive; feline; protein; immune response; antigen; tolerance; mammal; Dermatophagoides; Felis; Ambrosia; Lolium; Canis; Cryptcomeria; Alternaria; Alder; Betula; Quercus; Olea; Artemesia; Plantago; Parietaria; Blattella; Apis; Periplaneta; autoantigen; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the
                                                                                        Sequence
                                                                                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                                                     WPI; 1993-320744/40
                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                     30-SEP-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9319178-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human T cell reactive feline protein fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
21-APR-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15
                                                                                                                                                                                                                                                                                                                                             (IMMU-) IMMUNOLOGIC PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention.
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                                            Similarity
5; Conser
                                                                                         16
                                                                                                                                                                                                                  Fig 3; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                     DKQNYLALVREL 12
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                                                                                                                                                                                                                                                                                                                      Garman
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                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
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92US-00884718.
93US-00006116.
                                                                                                                                                                                                                                                                                                                                                                                                                 93WO-US002462
                                                                                                                                                                                                                                                                                                                      ВD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43.8%;
                                                      43.8%;
41.7%;
                                                                                                                                                                                                                                                                                                                        Greenstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16
                                          Score 28; DB Pred. No. 5.5e 5; Mismatches
                                                                                                                                                                                                                                                 subcutaneously, for t
                                            5
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Pred. No. 5.1e+02;
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                                                                                                                                                                                                                                                                                                                      JĽ,
                                                      DB 2; I
5.5e+02;
                                                                                                                                                                                                                                                                                                                      Kuo
                                                                                                                                                                                                                                                                                                                      Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 15;
                                                                Length 16;
                                                                                                                                                                                                                                                 e one or more epitope(s)
treating sensitivity to
                                                                                                                                                                                                                                                                                                                        Briner
                                                                                                                                              which involve an immune ay be used to induce Ambrosia, Lolium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                             Indels
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um; Canis;
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                                            Gaps
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AAB28946 standard; peptide;

16

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RESULT 20
AAY51484
ID AAY51484
ID AAY51484;
XX
AC AAY51484;
XX
AC AAY51484;
XX

DT 22-MAY-2000 (first entry)
XX

DE Human TRFP derived peptide C.
XX

T-cell reactive feline protein; T
KW down regulation; inmune response;
KW sensitivity; cat protein allergen
XX

OS Homo sapiens.
XX

PD 01-FEB-2000.
XX

PF 02-SEP-1994; 94US-00300928.
XX

PF 02-SEP-1991; 91US-00662276.
PR 28-FEB-1991; 91US-00662276.
PR 28-FEB-1991; 91US-00867311.
PR 25-MAY-1992; 92US-00887311.
PR 15-JAN-1993; 93US-00887311.
PR 15-JAN-1993; 93US-00066116.
XX

PF 03-NOV-1989; 92US-00887711.
PR 15-JAN-1993; 93US-00066116.
XX

PF 15-JAN-1993; 93US-00066116.
XX

PF 25-MAY-1992; PRARM CORP.
XX

PF 25-MAY-1993; PR 25-MAY
RESULT 21
AAB28946
ID AAB2
                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel peptide (I) of human T cell reactive feline protein (hTRFP) having at least one T cell epitope recognized by a T cell receptor specific for the human T cell reactive feline protein, the peptide consisting of at least 7-30 amino acids, and having an amino acid sequence derived from an amino acid sequence comprising 94, 96, 97, 109, or 111 residues, given in the specification. The peptides down regulate the immune response to the allergen. The peptides have reduced immunoglobulin B binding and reduce T cell responsiveness. The peptide (I) is useful in compositions for treating sensitivity to a cat protein allergen in a subject. This sequence represents the human TRFP derived peptide used in trhe method of the invetnion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptides of human T cell reactive feline protein for treating sensitivity to cat protein allergens comprise at least one T cell epitope recognized by a T cell receptor specific for the human T cell reactive feline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  T-cell reactive feline protein; TRFP; T cell epitope; T cel down regulation; immune response; allergen; immunoglobulin
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                                                                                                                                                                                                                                                                                Similarity
5; Conserv
                                                                                                                                                               DKENALSLLDKI
                                                                                                                                                                                                                      DKQNYLALVREL
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                                                                                                                                                                                                                                                                                                             43.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kuo
                                                                                                                                                                                                                                                                                       5
                                                                                                                                                                                                                                                                                                                 Score 28; I
Pred. No. 5.
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                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                          DB 3;
5.5e+02;
2;
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                                                                                                                                                                                                                                                                                                                                              Length 16;
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RESULT 22
AAY90112
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Best Local S
Matches
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15-MAY-1992;
15-JAN-1993;
02-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-NOV-1989;
28-FEB-1991;
13-DEC-1991;
                                                                                                                                                                                                                                                                                                The present invention relates to the detection of sensitivity to a cat protein allergen by combining a blood sample from a subject with a peptide of human T cell reactive feline protein (hTRFP). This method and the hTRFP peptides are useful for diagnosing, preventing and treating cat allergies by reducing or abolishing an individual's allergic response to a cat allergen. DNA encoding the TRFP may be used as probes to locate equivalent sequences present in other species. These may further be used to study the mechanism of immunotherapy of cat allergy, and to design modified derivatives, analogues or functional equivalents useful in immunotherapy. The present sequence was used in the invention
                           Cat; TRFP; human T-cell reactive feline protein; cat protein allergen; house dust; Fel d I; cat allergy; Felis domesticus sensitivity; therapy; diagnosis; goat; sheep; horse; rabbit; dog.
                                                                                         12-SEP-2003
13-JUL-2000
                                                                                                                                           AAY90112 standard; peptide; 16 AA
                                                                                                                                                                                                                                                                                Sequence 16 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting, preventing and treating sensitivity to cat protein allergen comprises combining a biological sample with a human T cell reactive feline protein and determining the extent of binding that occurs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gefter ML,
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US6120769-A
         Felis catus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide C derived from T cell reactive feline protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IMMU-) IMMULOGIC PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-APR-1995;
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                                                                   TRFP derived peptide, peptide C
                                                                                                                                                                                                                  μ.
                                                                                                                                                                                                                                      Similarity
5; Conserv
                                                                                                                                                                                                        DKQNYLALVREL 12
                                                                                                                                                                                               DKENALSLLDKI 14
                                                                                                                                                                                                                                                                                                                                                                                                                    Fig 17; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Garman RD,
                                                                                                                                                                                                                                        Conservative
                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92US-00857311.
92US-00884718.
93US-00006116.
94US-00300928.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human T cell reactive feline protein; hTRFP; immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       91US-00662276.
91US-00807529.
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                                                                                                                                                                                                                                                43.8%;
41.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Greenstein
                                                                                                                                                                                                                                    Score 28; DB 3; Le
Pred. No. 5.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bond JF
                                                                                                                                                                                                                                                         Length 16;
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03-NOV-1989;
28-FEB-1991;
13-DEC-1991;
                                                                                                                                                       This sequence is a peptide derived from the human T cell reactive feline protein (TRFP) of the invention. The protein is a cat protein allergen, and was isolated from a vacuum bag extract obtained by affinity purification of house dust collected from several homes with cats. TRFP is composed of two covalently linked peptide chains, and is also referred to as Fel d I. TRFP and its peptides are useful for reducing or preventing the adverse effects that exposure to cat allergens normally has on cat allergic individuals (i.e. to desensitise individuals to cat allergens or block the effect of the allergens). TRFP also used in methods of diagnosing sensitivity to Felis domesticus in an individual. DNA sequences present in other species, e.g. goat, sheep, horse, rabbit and dog, that may be useful in diagnostic and/or therapeutic applications. (Updated on 12-SEP-2003 to standardise OS field)
                                                                                                             Sequence 16 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New human T cell reactive feline protein, allergic individuals to cat allergens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-181812/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JAN-1993;
02-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-APR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (IMMU-) IMMULOGIC PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9; Fig 17; 108pp; English
Similarity 41. 5; Conservative
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91US-00662276.
91US-00807529.
92US-00857311.
92US-00884718.
93US-00066116.
94US-00300928.
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                            43.8%;
41.7%;
υ
••
Score 28; DB 3; Le
Pred. No. 5.5e+02;
5; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful for desensitizing
                                                     Length 16
  Indels
  0
  Gaps
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Claim

RESULT 23 AAY87687 밁 á Query Match Best Local : Matches 03-NOV-1989; 27-APR-1995; US6048962-A. Felis sp. T-cell reactive antiallergic; T Feline human TRFP peptide C. AAY87687 standard; protein; 16 AA 11-APR-2000. 22-AUG-2000 AAY87687; w 1 DKONYLALVREL 12 (first entry) 89US-00431565 9508-00430014 feline cell st ne protein; '
stimulator; TRFP; Fel TRFP; Fel d diagnostic; H immunotherapy.

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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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13-DEC-1991;
25-MAR-1992;
15-MAY-1992;
15-JAN-1993;
02-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptides cause sensitivity so as to selectively use them to desensitize a cat sensitive individual. Purified TRFP is also useful for studying the mechanism of immunotherapy of cat allergy and to design modified derivatives, analogs or functional equivalents that are more useful in immunotherapy against cat allergy. DNA sequences encoding TRFP are useful as probes to locate equivalent sequences present in other species (goats, sheep, dogs, rabbits or horses) that may be useful in diagnostics and/or therappeutics. Fully defined and characterized TRFP provides complete and a very simple desensitization therapy. This sequence represents a human T cell reactive feline protein (also known as Fel d I) derived peptide C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel naturally occurring cat protein allergen (I), human T cell reactive feline protein (TRFP), comprising two different covalently linked peptide chains with a molecular weight of 20 kD, 40 kD or 130 kD under non-reducing conditions and 5 kD or 10-18 kD under reducing conditions. The products of the invention have antiallergic activity and act as human T cell stimulators. TRFP is useful for reducing or preventing the adverse effects of cat allergens on cat allergic individuals and in ex vivo diagnostic tests to determine which
                 WO2003046204-A2
                                                                                                    murine;
                                                                                                                    anti-cryptic collagen; HUIV26; variable region light chain; mouse;
                                                                                                                                   grafted antibody; complementarity determining region; CDR; light CDR; heavy CDR; cryptic collagen epitope; solid tumour; new blood vessel growth; angiogenesis; tumour growth; cytostatic; collagen agonist; collagen antagonist; cancer metastasis;
                                                                                                                                                                                                                        Mouse HUIV26 antibody mutant light chain CDR amino acid sequence SeqID76
                                                                                                                                                                                                                                                           29-JAN-2004
                                                                                                                                                                                                                                                                                             ADD94191;
                                                                                                                                                                                                                                                                                                                           ADD94191 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   which is described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human T cell reactive feline protein useful for reducing or abolishing individual's allergic response to cat allergen comprising different covalently linked peptide chains.
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ein JL, Griffith IJ,
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92US-00857311.
92US-00884718.
93US-00006116.
94US-00300928.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New cryptic collagen antibody with one or medetermining regions, useful for diagnosing associated with angiogenesis, tumor growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Recombinant Fel d 1; rFel d 1; Felis domesticus allergen Fel d 1 chain 1; Fel d 1 chain 2; fusion product; cat all antiallergic; domestic cat; mutant; mutein; tryptic fragm
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71.4%;
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Groenlund H,

Van Hage- Hamsten

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Search completed: January 26, 2006, 07:57:55 Job time : 48.1552 secs
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                                                                                                                                                                                                                                                                                                                      The invention relates to a recombinant Felis domesticus allergen 1 (rFel Cd 1) fusion product comprising a Fel d 1 chain 2 and a CC linker, e.g. a carbon-nitrogen bond or a short peptide linker, which CC links the N-terminal amino acid of one chain to the C-terminal amino acid CC of the other chain. Also disclosed are (i) a homodimer consisting of two non-covalently associated fusion products, (ii) a DNA sequence encoding CC the fusion product, (iii) an expression vector having the DNA sequence CC inserted into it in an operable form, (iv) a host cell transformed with CC the expression vector, (v) a pharmaceutical composition comprising an CC immunotherapeutical manunt of the fusion product and/or the homodimer and a pharmaceutical carrier, excipient or diluent, (vi) a kit, for CC diagnosing cat allergy, comprising the fusion product and/or the homodimer and (ix) a process for preparing a recombinant Fel d l fusion product comprises a fully defined sequence of CC and (ix) a process for proparing a recombinant Fel d l fusion product comprises a fully defined sequence of CC at allergy. The composition, kite, and method are useful for diagnosing, treating or preventing cat allergy. This sequence represents a tryptic CC fragment of the rFel d 1 (2+1) fusion product.
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                                                                                                                                                                                                                                                                                   Sequence 17 AA;
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US-08-430-944D-22
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US-08-430-928C-22
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1851, Ap
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ALIGNMENTS

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APPLICANT: MCCALI, Catherine A.
APPLICANT: HURLEY, Shirley Wu
APPLICANT: HURLEY, Shirley Wu
APPLICANT: HURLEY, Shirley Ru
APPLICANT: HURLEY, Shirley Ru
APPLICANT: HURLEY, STIC R.
TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PRO;
TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PRO;
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: AL-2-C3
CURRENT APPLICATION NUMBER: US/09/292,225
CURRENT FILING DATE: 1999-04-15
EARLIER APPLICATION NUMBER: 60/098,909
EARLIER FILING DATE: 1998-05-13
EARLIER APPLICATION NUMBER: 60/085,295
EARLIER APPLICATION NUMBER: 60/08,565
EARLIER APPLICATION NUMBER: 60/08,565
EARLIER APPLICATION NUMBER: 09/08,565
EARLIER APPLICATION NUMBER: 09/08,565
EARLIER APPLICATION NUMBER: 09/08,565
EARLIER APPLICATION NUMBER: 09/08,565
EARLIER FILING DATE: 1998-04-17
NUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BOOTS, ANNA M.H.

APPLICANT: VERHEIJDEN, GILBERTUS F.M.

TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR

TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY

FILE REFERENCE: 0/96198 US

CURRENT APPLICATION NUMBER: US/09/171,705

CURRENT FILING DATE: 1999-02-09

NUMBER OF SEQ ID NOS: 78

SOFTWARE: PATENTIN Ver. 2.0

SEQ ID NO 21
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US-09-292-225-7
                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: DERIVED FROM ; OTHER INFORMATION: SEQUENCE OF HUMAN CARTILAGE (HC) -39 PROTEIN US-09-171-705-21
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                                                                                        Query Match
Best Local S
Matches 5
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Patent No. 6455686
GENERAL INFORMATION:
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Best Local
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                                                                                                                                                                                                                                                           LENGTH: 16
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                        Similarity 5; Conserv
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                                            DKQNYLALVREL 12
                                                                                           Conservative
                                                                                                              54.7%;
41.7%;
                                                                                        Score 35; DB 2;
Pred. No. 10;
5; Mismatches
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                                                                                                                                       Length 16;
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APPLICANT: FUJITA, Tomoko
APPLICANT: MIYAKE, Kazue
APPLICANT: MIYAKE, Kazue
APPLICANT: MIYAKE, Kazue
APPLICANT: MSADA, Kiyozo
APPLICANT: KATO, Ikunoshin
TITLE OF INVENTION: DNA POLYMERASE-RELATED FAV
FILE REFERENCE: 1422-408PCT
CURRENT APPLICATION NUMBER: US/09/446,504
CURRENT FILING DATE: 1999-12-23
PRIOR FILING DATE: 1998-06-24
PRIOR FILING DATE: 1998-06-24
PRIOR APPLICATION NUMBER: UF 9-187496
PRIOR APPLICATION NUMBER: JP 9-20692
PRIOR FILING DATE: 1997-06-26
PRIOR APPLICATION NUMBER: JP 9-320692
PRIOR FILING DATE: 1997-11-27
PRIOR FILING DATE: 1997-11-27
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Best Local Similarity
Similarity
Conserve
                        Query Match
Best Local Similarity
Whiches 6; Conserve
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; OTHER INFORMATION: Description of Artificial Sequence: DERIVED FROM
; OTHER INFORMATION: SEQUENCE OF HUMAN CARTILAGE (HC) -39 PROTEIN
US-09-657-757-21
                                                                                                                             LENGTH: 20
TYPE: PRT
CORGANISM: Pyrococcus furiosus
US-09-446-504-7
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APPLICANT: BOOTS, ANNA M.H.
APPLICANT: BOOTS, ANNA M.H.
APPLICANT: VERHELIDEN, GILBERTUS F.M.
APPLICANT: VERHELIDEN, GILBERTUS F.M.
TITLE OF INVENTION: INMUNOSUPPRESSIVE THERAPY
FILE REFERENCE: 0/96198 US
CURRENT APPLICATION NUMBER: US/09/657,757
CURRENT APPLICATION NUMBER: US/09/171,705
PRIOR APPLICATION NUMBER: US/09/171,705
PRIOR FILING DATE: 1998-10-23
                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin V
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 78
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21, Appl:
Patent No. 688182
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Patent No. 6218150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: UEMORI, Takashi
APPLICANT: SATO, Yoshimi
APPLICANT: FUJITA, Tomoko
APPLICANT: MIYAKE, Kazue
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ORGANISM: Artificial Sequence
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DKQNYLALVRE 11
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milarity 41.7%;
Conservative
                                           Conservative
                                                               46.9%;
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Pred. No. 10;
                                                               Score 30; DB
Pred. No. 97;
                                              Mismatches
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; ORGANISM: Pyrococcus furiosus
US-09-712-266-7
                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 22
LENGTH: 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. SEQ ID NO 7
                                                                            Matches
                                                                                                              Query Match
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Best Local (
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CURRENT APPLICATION NUMBER: US/09/171,705
CURRENT FILING DATE: 1999-02-09
NUMBER OF SEQ ID NOS: 78
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: BOOTS, ANNA M.H.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR USE IN ANTIGEN SPECIFIC
TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 7, Applicat
Patent No. 6333158
GENERAL INFORMATION:
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CURRENT FILING DATE: 2000-11-15
PRIOR APPLICATION NUMBER: US 09/446,504
PRIOR FILING DATE: 1999-12-23
PRIOR FPLICATION NUMBER: PCT/JP98/02845
PRIOR FILING DATE: 1998-06-24
PRIOR FILING DATE: 1998-06-24
PRIOR FILING DATE: 1998-06-24
PRIOR APPLICATION NUMBER: JP 9-187496
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APPLICANT: ASADA, Kiyozo
APPLICANT: KATO, Ikunoshin
TITLE OF INVENTION: DNA POLYMERASE-RELATED FACTORS
FILE REFERENCE: 1422-408PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: UEMORI, Takash
APPLICANT: SATO, Yoshimi
                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                  TYPE: PRT
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                                                                                              Local
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                                                                                              Similarity
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QHFTTLIKEMK 11
                                  QNYLALVRELK 13
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MIYAKE, Kazue
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                                                                          Conservative
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                                                                                        45.38;
36.48;
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                                                                        Score 29; DB 2; Le
Pred. No. 1.2e+02;
5; Mismatches 2;
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Pred. No. 97;
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APPLICANT: BOOTS, ANNA M.H.
APPLICANT: VERHEIJDEN, GIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 601997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Pa
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        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/807,529
FILLING DATE: December 13, 1991
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: AMY E. MANDRAGOURAS
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: 002.6US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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PRIOR FILING DATE: 1998-10-23
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/300,928C
                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U
ZIP: 02145
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CTREET: 610 LINCOLN STREET
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(617)
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                                                                                                                                                                                                                                                                                      ASCII text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GEFTER, Malcolm L. et al.
/ENTION: PEPTIDES FOR HUMAN T CELL REACTIVE FELINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IMMULOGIC PHARMACEUTICAL CORPORATION
                                                                                                                                                                                                                                                                                                                                          Floppy disk
                                                                                                                                                                                                                          September 2, 1994
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36.4%;
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                                                    002.6US(IMI-044)
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Pred. No. 1.2e+02;
Predrehes 2;
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                                                                                                                           US-08-430-944D-22
                                                                                                                                                                                                                                                                                             SOFTMALE: BATCHING FLOW, Wersion #1.25
SOFTMARE: BATCHIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/8/430,944D
FILING DATE: 28-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/430,014
FILING DATE: 27-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/300,928
FILING DATE: 02-SEPT-1994
APTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IM-044DV2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6025162
GENERAL INFORMATION:
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Best Local (
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                                                              Matches
                                                                                                                                                                                                                                       TELEFAX: (617)742-421
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide FRAGMENT TYPE: internal
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                                                                                                                                          MOLECULE TYPE:
FRAGMENT TYPE:
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TELEPHONE: (617)742-4214
TELEPHONE: (617)742-4214
TELEPHONE: (617)742-4214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                       LENGTH:
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                                                             Similarity
5; Conserv
                                                                                                                                                                                       H: 16 amino acids amino acid
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DKENALSLLDKI 14
                            DKONYLALVREL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: LAHIVE & COCKFIELD, LLP
28 State Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bruce L. Rogers et al.
VENTION: A HUMAN T CELL REACTIVE FELINE PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                              Conservative
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                                                                                                                                                                           linear
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                                                                                                                                          peptide
internal
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                                                                             43.8%;
41.7%;
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                                                             Score 28; DB Pred. No. 1.7e 5; Mismatches
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Pred. No. 1.7e+02;
                                                                             DB 2;
1.7e+02;
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                                                                                          Length 16;
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                                                              Gaps
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RESULT 10

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; MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-430-014-22
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                                                                                                                                                                                                                                                                                                                   RESULT 11
                                                                                                                                                                                                   ; Patent No. 6120769
; GENERAL INFORMATION:
; APPLICANT: Bruce L. F
TITLE OF INVENTION: F
; TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22, Applicat Patent No. 6048962 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                      Sequence 22, Application US/08431184 Patent No. 6120769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617) 227-5941 INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                        NUMBER OF SEQUENCES: 103
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAWE: AMY E. MANDRAGOURAS
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: 002.6US (IMI-044)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: b..
CTTY: WALTHAM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
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                                                                                 COUNTRY:
                                                                                                                 CITY: Boston
                                                                                                                                   STREET:
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                                                                   02109
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                                                                                                    Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08430014
                                                                                                                                 28 State Street
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                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    43.8%;
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PEPTIDES FOR HUMAN T CELL REACTIVE FELINE
PROTEIN (TRFP)
                                                                                                                                                                                                                    Rogers et al.
A HUMAN T CELL REACTIVE FELINE PROTEIN
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Pred. No. 1.
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1.7e+02;
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JS-08-431-184-22
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                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/006,116

FILING DATE: 15-JAN-1993

APPLICATION NUMBER: US 07/884,718

FILING DATE: 15-MAY-1992

APPLICATION NUMBER: 07/857,311

FILING DATE: 25-MAR-1992
                                                                                                                                               SOFTWARE: ASCII TEXT CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT, FILING DATE: 1930325 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617)742-4214
[NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 27-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/300,928
FILING DATE: 02-SEPT-1994
ATTORNEY/AGENT IMPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy
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FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Gefter, Malcolm L. APPLICANT: Garman, Richard D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (617)227-7400
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                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII TEXT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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REFERENCE/DOCKET NUMBER: IMI-044DV3
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                                                                                                                                                                                                                                                                                                                                                                             CITY: Boston
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                                                                                                                                                                                                                                                                                                                                                                                                60 State Street
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Morville, Malcolm
VENTION: PEPTIDES USEFUL FOR TOLERIZATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kuo, Mei-chang
                                                                                                                                                                                                                                                                                                                                            USA
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internal
                                                                                                                                                                                                                                                                                 Floppy disk
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41.7%;
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Pred. No. 1.7e+02;
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; MOLECULE TYPE:
; FRAGMENT TYPE:
US-07-807-529A-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                           COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/807,529A
FILING DATE: 19911213
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/662,276
FILING DATE: 28-FEB-1991
APPLICATION NUMBER: US 07/431,565
FILING DATE: 03-NOV-1989
ATTORNEY/AGENT INFORMATION:
NAME: Channing Stack I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                          TELEPHONE: (617) 494-
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                           REGISTRATION NUMBER: 31,095
REFERENCE/DOCKET NUMBER: IPC-027/imi-015
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                         SEQUENCE CHARACTERISTICS:
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LENGTH: 16 amino acids
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FRAGMENT TYPE:
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                                                 LENGTH: 19 amino
TYPE: AMINO ACID
TOPOLOGY: 1 linear
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Local Similarity 41.7%;
nes 5; Conservative
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ZIP: 02139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cambridge
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                                                                                     19 amino acids
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/ENTION: RECOMBITOPE PEPTIDES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Greenstein, Julia L.
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Bond, Julian F.
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internal
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Pred. No. 1.
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Query Match Best Local Similarity

Score 28; DB 1; Pred. No. 2.1e+02;

Length 19;

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RESULT 15
US-08-430-944D-20
; Sequence 20, Application US/08430944D
"""APART NO. 6035162
"""APART NO. 6035162
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US-08-300-928C-20
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Patent No. 60132...
PARRAL INFORMATION:
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                                                               Patent No. 6025162
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Bruce L. Rogers et
TITLE OF INVENTION: A HUMAN T
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                Query Match
Best Local (
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TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 20:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/807,529
FILING DATE: December 13, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: AMY E. MANDRAGOURAS
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: 002.6US(IMI-044)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
 NUMBER OF SEQUENCES: 103
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/300,928C
FILING DATE: September 2, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
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FRAGMENT TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: VINERALTHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEULUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: UZIP: 02145
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                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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5; Conserve
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610 LINCOLN STREET
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28 State Street
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VENTION: PEPTIDES FOR HUMAN T CELL REACTIVE FELINE
                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     internal
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Pred. No. 2.
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CELL REACTIVE FELINE PROTEIN
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2.1e+02;
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US-08-430-014-20
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/430,944D
FILING DATE: 28-APR.1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/430,014
FILING DATE: 27-APR.1995
PRIOR APPLICATION DATA:
US 08/300,928
PRIOR APPLICATION NUMBER: US 08/300,928
APPLICATION NUMBER: US 08/300,928
APPLICATION INFORMATION:
NAME: AMY E. MANDERGOUTAS
REGISTRATION NUMBER: 36,207
                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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LENGTH: 19 amino acids
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
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TITLE OF INVENTION: PEPTIDES FOR HUMAN T CELL REACTIVE FELINE
TITLE OF INVENTION: PROTEIN (TRFP)
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMULOGIC PHARMACEUTICAL CORPORATION
                                                  CLASSIFICATION: PRIOR APPLICATION NUMBER:
                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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 ATTORNEY/AGENT INFORMATION:
                                                                                                     APPLICATION NUMBER: US/08/430,014 FILING DATE:
                                                                                                                                                                                                                                                                                  CITY: WALTHAM
STATE: Massacl
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REFERENCE/DOCKET NUMBER: IMI-044DV2
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                    CLASSIFICATION:
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41.7%;
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Pred. No. 2.1e+02;
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US-08-431-184-20
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Query Match 43.8%;
Best Local Similarity 41.7%;
Matches 5; Conservative
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Best Local Similarity 41.7%;
                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/431,184
FILING DATE: 28 APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/430,014
FILING DATE: 27-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/300,928
FILING DATE: 02-SEPT-1994
ATTORNEY/AGENT INFORMATION:
NAME: AMY E. MANDERSON ENGLISTRATION NUMBER: 36,207
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-044DV3
TELECOMMUNICATION INFORMATION:
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APPLICANT: Bruce
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                                                                                                                                                        TELEFAX: (617)742-4214
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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LENGTH: 19 amino acids
TYPE: amino acid
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPARAY. (---)
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CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bruce L.
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
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internal
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A HUMAN T CELL REACTIVE FELINE PROTEIN
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Score 28; DB 2; Le
Pred. No. 2.1e+02;
5; Mismatches 2;
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Pred. No. 2.1e+02;
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                                Length 19;
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RESULT 19
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PCT-US93-02462-10
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                                                             Sequence 14, Application PC/TUS9302462 GENERAL INFORMATION:
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Best Local Similarity 41.7%;
Matches 5; Conservative
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                       TOPOLOGY: 1i:
MOLECULE TYPE:
FRAGMENT TYPE:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 25-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: ASCII TEXT CURRENT APPLICATION DATA:
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ADDRESSEE: LAHIVE & COCKFIELD
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                                              APPLICANT:
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TITLE OF IN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/006,116
FILING DATE: 15-JAN-1993
APPLICATION NUMBER: US 07/884,718
FILING DATE: 15-MAY-1992
APPLICATION NUMBER: 07/857,311
                                                                                                                                                                                                                                                                                                                                                     TYPE: AMINO ACID
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OPERATING SYSTEM:
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STATE: MA
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                                                                                                                                                                                                                                                                                                                                                                     19 amino acids
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             Gefter, Malcolm L.
Garman, Richard D.
Greenstein, Julia L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Morville, Malcolm
VENTION: PEPTIDES USEFUL FOR TOLERIZATION
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Garman, Richard D.
Greenstein, Julia L.
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Briner, Thomas J.
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SYSTEM: PC-DOS/MS-DOS
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internal
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                                                                                                                                                                                                                                        Score 28; DB 4;
Pred. No. 2.1e+02;
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Best Local Similarity
"~~hes 5; Conserve
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                                                                                                                                                                                                                                            Patent No.
GENERAL IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/
FILING DATE: 15-JAN-1993
APPLICATION NUMBER: US 07/
FILING DATE: 15-MAY-1992
APPLICATION NUMBER: 07/857
FILING DATE: 25-WAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT,
FILING DATE: 1930325
CLASSIFICATION:
                                                                                                                                                                       APPLICANT:
APPLICANT:
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                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                         APPLICANT: Kuo, Mei-chang
APPLICANT: Morvile, Malcolm
                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
US 08/006,116
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COMPUTER READABLE FORM:
                                                                                                             TITLE OF INVENTION: RECOMBITOPE PEPTIDES NUMBER OF SEQUENCES: 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII TEXT
                                       STREET: One Kenderidge STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IPC-031PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
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            COUNTRY: U
ZIP: 02139
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                                                                                                                                                                                                                                               INFORMATION:
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                                                                                                                                                                                                                                                              7, Application US/07807529A 5547669
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               MA
USA
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AMINO ACID
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                                                                    One Kendall Square, Building 600
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Greenstein, Julia L.
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Bond, Julian F.
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                                                                                   IMMULOGIC PHARMACEUTICAL CORPORATION
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internal
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Pred. No. 2.2e+02;
5; Mismatches 2
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FILING DATE: 19911213
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/662,276
FILING DATE: 28-FEB-1991
APPLICATION NUMBER: US 07/431,565
                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-D
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/300
FILING DATE: September 2, 199
CLASSIFICATION: 435
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMULOGIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: GEFTER, TITLE OF INVENTION:
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LENGTH: 11 amino acids
TYPE: AMINO ACID
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07
FILING DATE: 03-NOV-1989
ATTORNEY/AGENT INFORMATION:
NAME: Channing, Stacey L.
REGISTRATION NUMBER: 31,0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-D
SOFTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
                             CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: AMY E. MANURAGOURAS
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: 002.6US(IMI-044)
TELECOMMUNICATION INFORMATION:
TELECHNNE: (617) 227-7400
                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/807,529
FILING DATE: December 13, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatil
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                 (617)
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SEQ ID NO: 27:
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55.6%;
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PEPTIDES FOR HUMAN T CELL REACTIVE FELINE
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                                                                                                                                                                                                                                                                                                                     PC-DOS/MS-DOS
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Pred. No. 1
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1.7e+02;
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SEQUENCE CHARACTERISTICS:

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RESULT 23
US-08-430-014-76
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                                                                                                                                  Query Match
Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                               TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          equence 76, Application US/08430944D atent No. 6025162 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/300,928
FILING DATE: 02-SEPT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragourae
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-044DV2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/430,944D FILING DATE: 28-APR-1995 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/430,014 FILING DATE: 27-APR-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
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TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                           LENGTH: 11 amino acids TYPE: amino acid TOPOLOGY: linear
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hes 5; Conserv
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STATE: Massa
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Application US/08430014
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                                                                                                                                     Conservative
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55.6%;
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A HUMAN T CELL REACTIVE FELINE PROTEIN
                                                                                                                                                 Score 27;
Pred. No. 1
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Pred. No. 1.
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1.7e+02;
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                                                                                                                                   1; Indels
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US-08-431-184-76
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GENERAL INFORMATION:
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Best Local Similarity 55.6%;
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TELEPAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 76:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Bruce
                  ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 103
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
STREET: 28 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Bruce L. Rogers et al.
TITLE OF INVENTION: A HUMAN T CELL REACTIVE FELINE PROTEIN
TITLE OF INVENTION:
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REFERENCE/DOCKET NUMBER: 00:
TELECOMMUNICATION INFORMATION:
CURRENT APPLICATION DATA
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NAME: AMY E. MANDRAGOURAS
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APPLICATION NUMBER:
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TYPE: Inear
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VENTION: PEPTIDES FOR HUMAN T CELL REACTIVE FELINE
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Sequence 9, Application US/09142885C

Patent No. 6737406

GENERAL INFORMATION:

APPLICANT: Kay, Anthony
APPLICANT: Larche, Mark

TITLE OF INVENTION: Cryptic Peptides and Method for Their Identification

FILE REFERENCE: 4028/75652

CURRENT APPLICATION NUMBER: US/09/142,885C

CURRENT FILING DATE: 1997-01-27

PRIOR APPLICATION NUMBER: PCT/GB97/00783

PRIOR FILING DATE: 1997-03-20

NUMBER OF SEQ ID NOS: 56

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 9

LENGTH: 16

TYPE: PRT

ORGANISM: Felis domesticus
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/430,014
PILING DATE: 27-APR-1995
PRIOR APPLICATION UMBER: US 08/300,928
PRIOR APPLICATION UMBER: US 08/300,928
PILING DATE: 02-SEPT-1994
ATTORNEY/AGENT INFORMATION:
NAME: AMY E. MANDERS 36,207
REFERENCE/DOCKET NUMBER: IMI-044DV3
TELECOMMUNICATION INFORMATION:
TELEPHAN: (617)742-4214
INFORMATION FOR SEQ ID NO: 76:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: protein
US-08-431-184-76
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Search completed: January 26, 2006, 08:06:59 Job time : 13.3822 secs
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Best Local Similarity 55.6%;
Matches 5; Conservative
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128.563 Million cell updates/sec
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2: 'cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: 'cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: 'cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: 'cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*

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US-09-995-529-76
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CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: US/09/292,225
PRIOR FILING DATE: 1999-04-15
PRIOR APPLICATION NUMBER: 60/098,909
PRIOR APPLICATION NUMBER: 60/085,295
PRIOR APPLICATION NUMBER: 60/085,295
PRIOR FILING DATE: 1998-09-02
PRIOR FILING DATE: 1998-09-02
PRIOR FILING DATE: 1998-09-02
PRIOR FILING DATE: 1998-09-03
                                                                  CURRENT APPLICATION NUMBER: US/09/971,309
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 09/446,504
PRIOR FILING DATE: 1999-12-23
PRIOR APPLICATION NUMBER: PCT/JP98/02845
PRIOR FILING DATE: 1998-06-24
PRIOR APPLICATION NUMBER: JP 9-187496
PRIOR PILING DATE: 1997-06-26
PRIOR PILING DATE: 1997-06-26
PRIOR APPLICATION NUMBER: JP 9-320692
PRIOR APPLICATION NUMBER: JP 9-320692
PRIOR FILING DATE: 1997-11-27
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SOFTWARE: PatentIn Ver. SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/09971309 Patent No. US20020106675A1
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APPLICANT: Hunter, Shirley Wu
APPLICANT: Weber, Eric R.
TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PROTEINS
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: AL-2-C3
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: UEMORI, Takash
APPLICANT: SATO, Yoshimi
APPLICANT: FUJITA, Tomok
                                                                                                                                                                                                                                                                                                                APPLICANT: ASADA, Kiyozo
APPLICANT: KATO, Ikunoshin
TITLE OF INVENTION: DNA POLYMERASE-RELATED FACTORS
FILE REFERENCE: 1422-0494P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. ?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/062,013 PRIOR FILING DATE: 1998-04-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/098,565 PRIOR FILING DATE: 1998-04-17
                                                   NUMBER OF SEQ ID NOS: 92
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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TYPE: PRT
ORGANISM: Dermatophagoides farinae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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MIYAKE, Kazue
                                                                                                                                                                                                                                                                                                                                                                                                              MUKAI, Hiroyuki
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Pred. No. 0.0002;
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; TYPE: PRT ; ORGANISM: Pyrococcus furiosus US-09-971-309-7
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ORGANISM: mycoplasma genitalium

FEATURE:

OTHER INFORMATION: Sequence located in MG017 at 15-24 and may interact with Sequence

OTHER INFORMATION: this patent.

US-09-573-822C-39
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US-09-573-822C-39
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                                                                                                    US-09-572-404B-2020
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US-09-572-404B-2020
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SOFWARE: ProtPatent version 1.0
SEQ ID NO 39
                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/572,404B
CURRENT FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 4203
SOFTWARE: ProtPatent version 1.0
SEQ ID NO 2020
LENGTH: 10
TYPE: PRT
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Best Local S
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Publication No. US20030199011A1
GENERAL INFORMATION:
APPLICANT: Proteom Ltd
TITLE OF INVENTION: Complementary peptide ligands generated from microbial genome sequence of invention in the patent
CURRENT APPLICATION NUMBER: US/09/573,822C
CURRENT FILING DATE: 2000-05-18
                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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                                    Matches
                                                  Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                            APPLICANT: Proteom Ltd
TITLE OF INVENTION: Complementary peptide ligands from the human genome
FILE REFERENCE: Human patent
                                                                                                            ORGANISM: Homo Sapiens
FEATURE:
OTHER INFORMATION: sequence located in SFRS8 OR SWAP
OTHER INFORMATION: Sequence 2019 in this patent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 10
                                                   Local Similarity
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 DKQNYLAL
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                                                   43.8%;
                                                   Score 28; DB 3;
Pred. No. 3.4e+02;
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 30; DB 3; Pred. No. 3.2e+02;
                                    Mismatches
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2.3e+02
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APPLICANT: Watkins, Jeffry D.
APPLICANT: Huse, William D.
APPLICANT: Tang, Ying
TITLE OF INVENTION: Humanized Collagen Antibodies and
TITLE OF INVENTION: Related Methods
FILE REFERENCE: P-1X 4976
CURRENT APPLICATION NUMBER: US/09/995,529
CURRENT FILING DATE: 2001-11-26
NUMBER OF SEQ ID NOS: 358
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Watkins, Jeffry D.
APPLICANT: Huse, William D.
APPLICANT: Tang, Ying
TITLE OF INVENTION: Humanized Collagen Antibodies and
TITLE OF INVENTION: Related Methods
FILE REFERENCE: P-1X 4976
CURRENT APPLICATION NUMBER: US/09/995,529
CURRENT FILING DATE: 2001-11-26
NUMBER OF SEQ ID NOS: 358
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 76
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                                                                                                                                        US-08-464-363-10
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                                                           Sequence 10, Application US/08464363
Publication No. US20030035815A1
GENERAL INFORMATION:
APPLICANT: Rogers, Bruce L.
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                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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Morgenstern, Jay
Bond, Julian F.
Garman, Richard D.
                                                                                                                                                                                                                                                                                                 Conservative
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71.4%;
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RESULT 8
US-10-463-113-10
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; FRAGMENT TYPE:
US-08-464-363-10
                                                                                                                                              Query Match 43.8%;
Best Local Similarity 41.7%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/807,529
FILING DATE: 13-DEC-1991
APPLICATION NUMBER: US 07/662,276
FILING DATE: 28-FEB-1991
APPLICATION NUMBER: US 07/431,565
FILING DATE: 03-MOV-1989
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-015CN
                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (617) 227-1
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/464,363
FILING DATE: 05-JUN-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Greenstein, Julia L.
APPLICANT: Kuo, Mei-chang
APPLICANT: MOTVILE, Malcolm
TITLE OF INVENTION: RECOMBITOPE PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII TEXT
                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                 TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Boston
STATE: MA
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                                                                                                                                                                                                                                                                                                                         19 amino acids
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                                                                                                                                                                                                                                                                                    linear
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                                                                                                                                                                                                                                              internal
                                                                                                                                                                                                                                                              peptide
                                                                                                                                                                    Score 28; DB 2;
Pred. No. 6.9e+02;
                                                                                                                                                                                    Length 19;
                                                                                                                                                  Indels
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Sequence 10, Application US/104 Publication No. US20040057959A1 GENERAL INFORMATION: COUNTRY: USA ZIP: 02109 COMPUTER READABLE FORM: CORRESPONDENCE ADDRESS:
ADDRESSEE: Lahive & Cockfield, LLP Morvile, Malcolm NUMBER OF SEQUENCES: CITY: Boston STATE: MA STREET: 28 State Street Application US/10463113 Morgenstern, Jay Bond, Julian F. Garman, Richard D. Greenstein, Julia L. Rogers, Bruce L. Kuo, Mei-chang

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RESULT 9
US-08-464-363-27
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICATION NUMBER: US/08/464,363
FILING DATE: 05-JUN-1995
PRIOR APPLICATION UNMBER: US 07/807,529
FILING DATE: 13-DEC-1991
APPLICATION NUMBER: US 07/662,276
                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                          APPLICANT: Greenstein, Julia L.
APPLICANT: Kuo, Mei-chang
APPLICANT: Morvile, Malcolm
TITLE OF INVENTION: RECOMBITOPE PEPTIDES
NUMBER OF SEQUENCES: 76
                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR DATE: 16-Jun-2003

PRIOR APPLICATION UNMBER: US 07/807,529

FILING DATE: 13-DEC-1991

APPLICATION NUMBER: US 07/662,276

FILING DATE: 28-FEB-1991

APPLICATION NUMBER: US 07/431,565

FILING DATE: 03-NOV-1989

ATTORNEY/AGNIT INFORMATION:

NAME: AMY E. MANDARSOURAB

REGISTRATION NUMBER: 36,207

REFERENCE/DOCKET NUMBER: 31-015CN

TELECHONE: (617) 742-4214
                                                                                                                                                                                                                    COUNTRY: UP ZIP: 02109
                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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SOPTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/463,113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO: 10:
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5; Conserva
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USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/08464363
                                                                                                                                                                                                                                                                                        60 State Street, Suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                Garman, Richard D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Morgenstern, Jay
Bond, Julian F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rogers, Bruce L.
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                                                                                                                                                                                                                                                                                                        Lahive & Cockfield
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43.8%;
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Pred. No. 6.9e+02;
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RESULT 10

US-10-463-113-27

; Sequence 27, Application US/104-

; Publication No. US20040057959A1

; GENERAL INFORMATION:
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Best Local Similarity
Marches 5; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: protein US-08-464-363-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 28-FEB-1991
APPLICATION NUMBER: US 07/431,565
FILING DATE: 03-NOV-1989
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-015CN
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                  TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 27:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                               PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/807,529
FILING DATE: 13-DEC-1991
APPLICATION NUMBER: US 07/662,276
FILING DATE: 28-FEB-1991
APPLICATION NUMBER: US 07/431,565
FILING DATE: 03-NOV-1989
ATTORNEY/AGENT INFORMATION:

NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
                                                                                                                                                                                                                                                                                                                                ZIP: 02109

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: ASCII TEXT

CURRENT APPLICATION NUMBER: US/10/463,113

FILING DATE: 16-Jun-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Lahive & Cockfield, LLP
STREET: 28 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Morvile, Malcolm
TITLE OF INVENTION: RECOMBITOPE PEPTIDES
NUMBER OF SEQUENCES: 76
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                          REFERENCE/DOCKET NUMBER: IMI-015CN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DKQNYLALV 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: MA
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                                                                       TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rogers, Bruce L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Garman, Richard D. Greenstein, Julia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Morgenstern, Jay
Bond, Julian F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kuo, Mei-chang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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Pred. No. 5.7e+02;
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Tark Charles

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TOPOLOGY: linear
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MOLECULE TYPE: protein
;
SEQUENCE DESCRIPTION: SEQ ID NO: 27:
US-10-463-113-27
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US-10-245-871-163
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 163
LENCTH.
                                                                                                                            PRIOR APPLICATION NUMBER: 09/396,813
PRIOR FILING DATE: 1999-09-14
NUMBER OF SEQ ID NOS: 905
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                               Sequence 163, Application US/10253286 Publication No. US20040058881A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 163, Application US/10245871 Publication No. US20030235594A1 GENERAL INFORMATION:
                                Query Match
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                                                                  ORGANISM: Felis domesticus -10-253-286-163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                             APPLICANT: XU, MINZHEN
TITLE OF INVENTION: 11-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2015
CURRENT APPLICATION NUMBER: US/10/253,286
CURRENT FILING DATE: 2003-01-13
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR PILING DATE: 2002-07-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Felis domesticus
-10-245-871-163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: XU, MINZHEN
TITLE OF INVENTION: 11-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2013
CURRENT APPLICATION NUMBER: US/10/245,871
CURRENT FILING DATE: 2003-01-09
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-07-17
                                                                                                                                                                                                                                                                                                              APPLICANT: HUMPHREYS, APPLICANT: XU, MINZH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/396,813 PRIOR FILING DATE: 1999-09-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: HUMPHREYS, ROBERT APPLICANT: XU, MINZHEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH: 16
Local Similarity 55. hes 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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                 42.2%;
55.6%;
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55.6%;
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                 Score 27; DB 4;
Pred. No. 8.6e+02;
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Pred. No. 8.6e+02;
3; Mismatches 1
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Pred. No.
 Mismatches
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 1; Indels
                                Length 16;
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Gaps
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US-10-805-650-104
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; TYPE: PRT
; ORGANISM: Felis catus
US-10-809-689-8
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US-10-702-400-104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-702-400-104
                                                          GENERAL INFORMATION
                                                                               Sequence 104, Application No.
                                                                                                                                                                                                                                                                            Query Match 42.2%;
Best Local Similarity 55.6%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/10809689 Publication No. US20040265342A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local
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APPLICANT: BORINS, Michael
TITLE OP INVENTION: DNA POlymerase Fusions and Uses Thereof
FILE REFERENCE: 25436/2382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/GB99/00080
PRIOR FILING DATE: 1999-01-11
PRIOR APPLICATION NUMBER: GB/9800445.0
PRIOR FILING DATE: 1998-01-09
PRIOR FILING DATE: 1998-01-09
PRIOR APPLICATION NUMBER: GB/9820474.6
PRIOR FILING DATE: 1998-09-21
NUMBER OF SEQ ID NOS: 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Methods and compositions for desensitisation FILE REFERENCE: 5538/1010 CURRENT APPLICATION NUMBER: US/10/809,689 CURRENT FILING DATE: 2004-03-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/702,400
CURRENT FILING DATE: 2003-11-05
NUMBER OF SEQ ID NOS: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Borns, Michael
TITLE OF INVENTION: DNA Polymerase Blends and Uses Thereof
FILE REFERENCE: 25436/2364
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 16
TYPE: PRT
ORGANISM: Methanopyrus kandleri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1
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                                                                            Application US/10805650 o. US20050048530A1
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Pred. No. 8.6e+02
                                                                                                                                                                                                                                                                                               Score 27; DB 5;
Pred. No. 8.6e+02;
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CURRENT APPLICATION NUMBER: US/10/805,650
CURRENT FILING DATE: 2004-03-19
PRIOR APPLICATION NUMBER: US 60/457,426
PRIOR FILING DATE: 2003-03-25
NUMBER OF SEQ ID NOS: 148
SOFTWARE: Patentin version 3.1
SEQ ID NO 104
LENGTH: 16
TYPE: PRT
                                                                                                                                                                                                                           US-10-044-692-63
Sequence 63, Application US/10044692
Publication No. US20030096344A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Methanopyrus kandleri US-10-805-650-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-994-595-46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 46, Application No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Microsoft Word 97
SEQ ID NO 46
LENGTH: 10
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local :
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APPLICANT: Suvarna, Kalavati
APPLICANT: Suvarna, Kalavati
APPLICANT: Bhattacherjee, Vasker
TITLE OF INVENTION: METHODS AND REAGENTS FOR DETECTING FUNGAL PATHOGENS IN
TITLE OF INVENTION: A BIOLOGICAL SAMPLE
FILLE REFERENCE: 96,247-A
CURRENT APPLICATION NUMBER: US/09/994,595
CURRENT FILING DATE: 2001-11-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT FILING DATE: 2001-11-27
PRIOR APPLICATION NUMBER: 08/650,809
PRIOR FILING DATE: 1997-05-20
NUMBER OF SEQ ID NOS: 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Polypeptide segment of ACVS_EMENI shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                Chapman, Karen D.
Morin, Gregg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT:
THERAPEUTIC METHODS
                                                                                                                                                                                                              APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                    N
                                                                                                                                                                                                                                                                                                                                                                                   5 YLALVRELK 13
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Similarity 66.7%;
6; Conservation
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Similarity 62.5%;
5; Conservative
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ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor
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                                                                                                                                                                                        Lingner, Joachim
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Pred. No. 8.6e
3; Mismatches
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Pred. No. 7.7e+02;
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RESULT 18
US-10-044-539-63
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                                                                                                                                                                                                       Sequence 63, Application US/10044539
Publication No. US20030100093A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 576-02
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 63:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/912,951
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,449
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/044,692
FILING DATE: 11-Jan-2002
CLASSIFFICATION: «Unknown»
PRIOR APPLICATION DATA:
                            TITLE OF
                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE DESCRIPTION: SEQ ID NO: 63:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Morin, Gralvin
Harley, Calvin
Andrews, William H.
Andrews, William H.
ANDENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND THERAPEUTIC METHODS
                                                                                                                                                                                      Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40.6%;
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
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SEQUENCE DESCRIPTION: SEQ ID NO: 63:
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REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 08/912,951
FILING DATE: <UNKnown
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APPLICATION NUMBER: US/10/044,539
FILING DATE: 11-Jan-2002
CLASSIFICATION: 435
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                         KQNYLALV 9
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OTHER INFORMATION: /note= "motif E peptide from Drosophila melanogaster TART non-LTR retrotransposable element reverse
                                                                                                                                                                                                                       KQSYLGVI 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: <Unknown>
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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CITY: San Francisco
STATE: California
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FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
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Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                               Mismatches
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KQSYLGVI 10

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US-10-325-810-63
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Best Local :
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FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
APPLICATION INFERR: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
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ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                          SEQUENCE DESCRIPTION: SEQ ID
                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide
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APPLICATION NUMBER: US/09/402,181
FILING DATE: 29-Sep-1997
APPLICATION NUMBER: US 08/724,643
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MEDIUM TYPE: Floppy disk
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Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                  Similarity
KQNYLALV 9
                                                                                                                                                                                                                   LOCATION: 1..11
OTHER INFORMATION: /note= "motif E peptide from Drosophila melanogaster TART non-LTR
                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/10/325,810 FILING DATE: 20-Dec-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
                                                                                                                                                                                                       retrotransposable element reverse
                                                                                                                                                                                                                                                                                                  NAME/KEY:
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/854,050 FILING DATE: 09-MAY-1997
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                                                                40.6%;
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                                      Score 26; DB 4;
Pred. No. 8.6e+02;
3; Mismatches 1
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                                                                                     Length 11;
                                             Indels
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RESULT 20
US-10-877-124-63
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                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 63:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                  NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,124
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TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                              FEATURE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
NAME/KEY: Peptide
LOCATION: 1.11
OTHER INFORMATION: /note= "motif E peptide from
Drosophila melanogaster TART non-LTR
                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/915,503 FILING DATE: 14-AUG-1997
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                 TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/911,312 FILING DATE: 14-AUG-1997
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                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                          LENGTH: 11 amino acids
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Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
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(415) 576-0300
                                                                                                              peptide
                                                                                                                                           linear
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Publication No. US2004
GENERAL INFORMATION:
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                                                                              APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/432,503
FILING DATE: 02-NOV-1999
APPLICATION NUMBER: 08/974,549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Cech,
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             NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/10/877,022 FILING DATE: 24-Jun-2004 CLASSIFICATION: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: <Unknown>
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Two Embarca
CITY: San Francisco
                                                                                                                                                FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/854,050 FILING DATE: 09-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
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                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/912,951 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                             APPLICATION NUMBER: US 08/915,503
                                                                                                                                                                                                                                                                          FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10877022
No. US20040247613A1
                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/911,312
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Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
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Pred. No.
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US-10-877-146-63
; Sequence 63, Application US/10877146
; Publication No. US20050013825A1
; GENERAL INFORMATION:
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 40.6%; Score 26; DB 5; Local Similarity 50.0%; Pred. No. 8.6e+02
                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/10/877,146
FILING DATE: 24-Jun-2004
CLASSIFICATION: <UNKnown>
                                                                                                                                                                           PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/432,503

FILING DATE: 02-NOV-1999

APPLICATION NUMBER: 08/974,549

FILING DATE: «Unknown»

APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997
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Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Peptide
LOCATION: 1..11
OTHER INFORMATION: /note= "motif E peptide from
Drosophila melanogaster TART non-LTR
retrotransposable element reverse
transcriptase"
SEQUENCE DESCRIPTION: SEQ ID NO: 63:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: Two Embarcadero CITY: San Francisco STATE: California
                                                                                   APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
                                          APPLICATION NUMBER: US 08/854,050 FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 576-0300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thomas R.
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RESULT 23
US-08-908-884-23
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APPLICATION NUMBER: US/08/908,884
FILLING DATE:
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/023,851
FILLING DATE: August 9,1996
APPLICATION NUMBER: 60/035,166
APPLICATION NUMBER: 60/035,166
FILLING DATE: January 10, 1997
APPLICATION NUMBER: 60/046,769
FILLING DATE: May 16, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Elbing, Karen L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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APPLICANT: Dong e
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHON: (415) 576-0200
INFORMATION FOR SEQ ID NO: 63:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                        OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Dong et al.
TITLE OF INVENTION: ACQUIRED RESISTANCE GENES AND USES THEREOF
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                ZIP:
                                                                                                                                                                                                                                                                                                                      COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Clark & Elbing of STREET: 176 Federal Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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LOCATION: 1.11
OTHER INFORMATION: /note= "motif E peptide from
Drosophila melanogaster TART non-LTR
retrotransposable element reverse
transcriptase"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
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REGISTRATION NUMBER: 36,429
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Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 11;
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TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 23: US-09-908-323-23
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Best Local Similarity
"---hes 5; Conserva
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acide
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                 NAME: Elbing, Karen L
REGISTRATION NUMBER: 35,238
REFERENCE/DOCKET NUMBER: 00786
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
TELEPAX: 617-428-7045
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23, Application US/09908323
Patent No. US20020073447A1
GENERAL INFORMATION:
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 35,238
REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION:
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                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/908,884
FILING DATE: <Unknown>
APPLICATION NUMBER: 60/035,166
FILING DATE: January 10, 1997
APPLICATION NUMBER: 60/046,769
FILING DATE: May 16, 1997
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/09/908,323
FILING DATE: 17-Jul-2001
CLASSIFICATION: -Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dong et al.
TITLE OF INVENTION: ACQUIRED RESISTANCE GENES AND USES THEREOF
NUMBER OF SEQUENCES: 28
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MEDIUM TYPE: Diskette
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ADDRESSEE: Clark & Elbing LLP
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Boston
STATE: MA
                                                                                                                              STRANDEDNESS: single
                                                                                                                                                TYPE: amino acids
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40.6%;
41.7%;
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Score 26; DB 3; Length 14; Pred. No. 1.1e+03;
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Search completed: January 26, 2006, 08:38:34
Job time : 44.25 secs
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Publication No. US20030068615A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 1242
SOFTWARE: PatentIn version 3.1
SEQ ID NO 684
LENGTH: 15
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                                                                                                                                                Matches
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Best Local Similarity 30.8%;
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CURRENT FILING DATE: 2002-02-27
PRIOR APPLICATION NUMBER: PCT/US00/23505
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/151,270
PRIOR FILING DATE: 1999-08-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: POLYPEPTIDES THAT BIND HIV GP120 AND RELATED NUCLEIC TITLE OF INVENTION: ACIDS, ANTIBODIES, COMPOSITIONS, AND METHODS OF USE FILE REFERENCE: 215875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SAXINGER,
                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: binding peptide
                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                     TYPE: PRT
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1: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*

8: /cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*
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1 DKQNYLALVRELK
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Maximum Match 100%
Listing first 100
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    GenCore version (c) 1993 - 2006
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  US-11-033-039-163
US-11-166-412-150
US-11-156-3-13-537-13
US-11-039-268-13
US-11-039-268-13
US-11-096-046-18
US-11-096-046-24
US-11-106-046-24
US-11-106-046-23
US-11-106-048-23
US-11-104-5-02-31
US-11-045-024-3576
US-11-045-024-3576
US-11-045-024-10581
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Sequence 163, App Sequence 13, Appl Sequence 13, Appl Sequence 18, Appl Sequence 24, Appl Sequence 24, Appl Sequence 27, Appl Sequence 27, Appl Sequence 31, Appl Sequence 6, Appl Sequence 6, Appl Sequence 1581, Appl Sequence 1585, Appl Sequence 10581, Appl Sequence 12381, Appl Sequence 12381, Appl Sequence 1381, Appl Sequence 3581, Appl Sequence 3581, Appl Sequence 3581, Appl Sequence 3582, Appl Sequence 358
                                                                                                                                                                                                                                                                               Description
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   US-11-045-024-10647
US-11-045-024-1240
US-11-045-024-1240
US-11-045-024-1240
US-11-045-024-12349
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112492, A

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1127, Appl

127, Appl

128, Appl

129, Appl

1296, Appl
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Sequence 163, Application US/11033039

Publication No. US20060002947A1

GENERAL INFORMATION:
APPLICANT: HUMPHREYS, ROBERT
APPLICANT: HUMPHREYS, ROBERT

FILE REFERENCE: REH-2017US01

FILE REFERENCE: REH-2017US01

CURRENT APPLICATION NUMBER: US/11/033,039

CURRENT FILING DATE: 2005-01-11

PRIOR APPLICATION NUMBER: 10/245,871

PRIOR APPLICATION NUMBER: 10/197,000

PRIOR FILING DATE: 2002-09-17

PRIOR APPLICATION NUMBER: 09/396,813

PRIOR APPLICATION NUMBER: 09/396,813

PRIOR FILING DATE: 1999-09-14
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US-11-166-412-150
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                                                                                                                                ; OTHER INFORMATION: Protein domain fragment US-11-166-412-150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: PRT; ORGANISM: Felis catus US-11-033-039-163
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US-11-033-039-163
                        Query Match
Best Local Similarity bb...
Conservative
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Publication No. US20060014231
GENERAL INFORMATION:
APPLICANT: Van Rompaey, Luc
APPLICANT: Tomme, Peter H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 163
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                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Tomme, Peter H. M.
TITLE OF INVENTION: Methods and Compositions To Promote Bone Homeostasis
FILE REFERENCE: PZ7, 927-D USA
CURRENT APPLICATION NUMBER: US/11/166,412
CURRENT FILING DATE: 2005-06-24
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SOFTWARE: PatentIn version 3.3
EQ ID NO 150
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SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/630,449
PRIOR EILING DATE: 2004-11-23
PRIOR APPLICATION NUMBER: 60/673,206
PRIOR FILING DATE: 2005-04-20
                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/582,704 PRIOR FILING DATE: 2004-06-24
                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                    ENGTH: 19
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Local Similarity 55.6%;
nes 5; Conservative
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                  5 YLALVRELK 13
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Pred. No. 19;
                                                                         Score 26; DB 7; Length 19; Pred. No. 35;
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Sequence 22, Appl
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CURRENT FILING DATE: 2005-01-19
PRIOR APPLICATION NUMBER: 09/620,840
PRIOR FILING DATE: 2000-07-21
PRIOR APPLICATION NUMBER: DIV of 09/620,840
PRIOR FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin version 3.3
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; LENGTH: 7
; TYPE: PRT
; ORGANISM: frog
US-10-753-537-13
                                                                                                                                                                                         ; OTHER INFORMATION: fragment having properties substantially similar to that of ; OTHER INFORMATION: leucine-based motif US-11-039-268-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-753-537-13
; Sequence 13, Application US/10753537
; Publication No. US20050260230A1
; GENERAL INFORMATION:
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US-11-039-268-13
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PRIOR FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin version 3.1
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13, Application US/11039268 Publication No. US20050276820A1 GENERAL INFORMATION:
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Best Local Similarity
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Herrington, Todd
APPLICANT: Aoki, Kei
TITLE OF INVENTION: LEUCINE-BASED MOTIF AND CLOSTRIDIAL NEUROTOXINS
FILE REFERENCE: ALLEO014-106 (17355DIV1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: STEWARD,
APPLICANT: FERNANDI
APPLICANT: HERRING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Steward, Lance APPLICANT: Herrington, To APPLICANT: Aoki, Kei
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CURRENT FILING DATE: 2004-01-08
PRIOR APPLICATION NUMBER: US/09/910,346C
PRIOR FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Leucine-based motif and clostridial neurotoxins FILE REFERENCE: D-2885CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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HERRINGTON, TODD
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                                                                                                                Score 25; DB 7; Le
Pred. No. 5.8e+04;
0; Mismatches 1;
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Pred. No. 5.8e+04;
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CURRENT APPLICATION NUMBER: US/10/981,356A
CURRENT FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: US 60/520,398
PRIOR FILING DATE: 2003-11-13
PRIOR APPLICATION NUMBER: US 60/557,951
PRIOR FILING DATE: 2004-03-31
NUMBER OF SEQ ID NOS: 45
SEQ ID NO 18
LENGTH: 17
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                                                                                                                                        RESULT 7
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; OTHER INFORMATION: Sequence is synthesized US-10-981-356A-18
                                                                                                                        US-11-096-046-18
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                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 45
SEQ ID NO 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24, Application US/10981356A Publication No. US20060015952A1 GENERAL INFORMATION:
APPLICANT: FILVAROFF, ELLEN H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18, Application US/10981356A Publication No. US20060015952A1 GENERAL INFORMATION:
                                                                     GENERAL
                                                                                     Sequence 18, Application US/11096046 Publication No. US20050276802A1
                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SCREENING ASSAYS AND METHODS OF TUMOR TREATMENT FILE REFERENCE: P2068R1
                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/557,951 PRIOR FILING DATE: 2004-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/981,356A CURRENT FILING DATE: 2004-11-04
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TITLE OF INVENTION: SCREENING ASSAYS AND METHODS OF TUMOR TREATMENT
FILE REFERENCE: P2068R1
                   APPLICANT:
                                                     APPLICANT: ADAMS,
                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                      INFORMATION:
                                                                                                                                                                                          11 NOKNYLA 17
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FERRARA, NAPOLEONE
FILVAROFF, ELLEN H.
MAO, WEIGUANG
                                                                                                                                                                                                                                                            Conservative
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                                                     CAMELLIA W.
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57.1%;
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57.1%;
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; Pred. No. 73;
3; Mismatches
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Pred. No. 73;
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US-11-105-708-23
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                                                                                                                                                                                    Sequence 23, Applica Publication No. US20 GENERAL INFORMATION:
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SEQ ID NO 18
LENGTH: 17
TYPE: PRT
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Best Local Similarity 57.1%;
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Best Local Similarity
APPLICANT: Pernasetti, Flavia
APPLICANT: Fréimark, Bruce
APPLICANT: Fréimark, Bruce
APPLICANT: Van Bpps, Dennis
APPLICANT: Brooks, Peter C
TITLE OF INVENTION: Method and Composition for Angiogenesis Inhibition
FILE REFERENCE: 30797-704.501
CURRENT APPLICATION NUMBER: US/11/105,708
CURRENT FILING DATE: 2005-04-13
PRIOR APPLICATION NUMBER: 09/478,977
PRIOR FILING DATE: 2000-01-06
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CURRENT FILING DATE: 2005-03-31
PRIOR APPLICATION NUMBER: US 60/558,290
PRIOR FILING DATE: 2004-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MAO, WEIGUANG
APPLICANT: PRESTA, LEONARD G.
APPLICANT: TEJADA, MAX L.
TITLE OF INVENTION: Humanized Anti-TGF-Beta Antibodies
FILE REFERENCE: P1954R1US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: TEJADA, MAX L.
TITLE OF INVENTION: Humanized Anti-TGF-Beta Antibodies
FILE REFERENCE: P1954R1US
CURRENT APPLICATION NUMBER: US/11/096,046
CURRENT FILING DATE: 2005-03-31
PRIOR APPLICATION NUMBER: US 60/558,290
PRIOR FILING DATE: 2004-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION:
                                                                                                                                                                                                                                                                                                                    11 NOKNYLA 17
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                                                                                                                                                                                                     Application US/11105708 No. US20050281821A1
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Pred. No. 73;
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PRIOR APPLICATION NUMBER: 60/152,496

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APPLICANT: PATEL, Pratiksha

APPLICANT: HALL, Andrea

APPLICANT: HALL, Andrea

FITTE OF INVENTION: MONOCLONAL ANTIBODIES TO THE CLFA PROTEIN . . . .

FILE REFERENCE: P07069US04/BAS

CURRENT APPLICATION NUMBER: US/11/136,559

CURRENT FILLING DATE: 2005-05-25

PRIOR APPLICATION NUMBER: US/10/056,052

PRIOR FILING DATE: 2002-01-18

PRIOR APPLICATION NUMBER: 60/308,116

PRIOR APPLICATION NUMBER: 60/298,413

PRIOR APPLICATION NUMBER: 60/298,413

PRIOR APPLICATION NUMBER: 60/298,413

PRIOR APPLICATION NUMBER: 60/274,611

PRIOR APPLICATION NUMBER: 60/264,072

PRIOR APPLICATION NUMBER: 2001-03-26

PRIOR APPLICATION NUMBER: 2001-03-26

PRIOR APPLICATION NUMBER: 2001-03-26

PRIOR APPLICATION NUMBER: 2001-03-26
RESULT 11
US-11-021-305-31
; Sequence 31, Application US/11021305
; Publication No. US20050282733A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                         ; TYPE: PRT ; ORGANISM: Staphylococcus aureus US-11-136-559-27
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US-11-136-559-27
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APPLICANT: PATTI, J
APPLICANT: HUTCHIN
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Publication No.
                                                                                                                                                                                                                           Matches
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Best Local Similarity 57.1%;
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PRIOR FILING DATE: 1999-01-06
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.2
SEQ ID NO 23
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1999-09-02
PRIOR APPLICATION NUMBER: 60/143,534
PRIOR FILING DATE: 1999-09-02
PRIOR APPLICATION NUMBER: 60/114,878
PRIOR FILING DATE: 1999-01-06
PRIOR FILING DATE: 1999-01-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Synthetic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                            PatentIn version 3.1
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DOMANSKI, Paul
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INS, Jeff T
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Pred. No. 73;
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Query Match
Best Local Similarity
Thes 5; Conserv
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US-11-041-893-201
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                                                                                                                                                                                                                                         Sequence 6, Application US/10494781 Publication No. US20050244895A1 GENERAL INFORMATION:
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Best Local Similarity 30...
Thes 4; Conservative
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 201, Application US/11041893 Publication No. US20060002941A1 GENERAL INFORMATION:
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SEQ ID NO 31
LENGTH: 12
TYPE: PRT
      APPLICANT: King's College London
TITLE OF INVENTION: Diagnosis of Demyelinating or Spongiform Disease
TITLE REPERENCE: IT/KE/N13246
CURRENT APPLICATION NUMBER: US/10/494,781
CURRENT FILING DATE: 2004-05-06
PRIOR APPLICATION NUMBER: GB 0127000.8
PRIOR PILING DATE: 2001-11-09
PRIOR FILING DATE: 2001-11-09
PRIOR FILING DATE: 2002-02-04
NUMBER OF SEQ ID NOS: 28
PRIOR FILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 28
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TITLE OF INVENTION: COMPOSITIONS COMPRISING IMMUNE RESPONSE
TITLE OF INVENTION: ALTERING AGENTS AND METHODS OF USE
FILE REFERENCE: 100123.401
CURRENT APPLICATION NUMBER: US/11/041,893
CURRENT FILING DATE: 2005-01-24
PRIOR APPLICATION NUMBER: US 60/616,855
PRIOR APPLICATION NUMBER: US 60/538,713
PRIOR PILING DATE: 2004-01-23
PRIOR FILING DATE: 2004-01-23
PRIOR FILING DATE: 2004-01-23
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APPLICANT: Hutley, Louise J
TITLE OF INVENTION: Differentiation-modulating agents and uses therefor
FILE REFERENCE: DAVI169.001CP1
FILE REFERENCE: DAVI169.001CP1
CURRENT APPLICATION NUMBER: US/11/021,305
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR FILING DATE: 2002-06-27
PRIOR FILING DATE: 2002-06-27
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SOFTWARE: PatentIn version 3.1
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TYPE: PRT
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62.5%;
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Pred. No. 1.4e+02;
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APPLICANT: Epimmune Inc.

APPLICANT: Epimmune Inc.

TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency

TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions

FILE REFERENCE: 2060.0040007

CURRENT APPLICATION NUMBER: US/11/045,024

CURRENT FILING DATE: 2005-01-28

PRIOR APPLICATION NUMBER: US 09/412,863

PRIOR APPLICATION NUMBER: US 09/027,146

PRIOR APPLICATION NUMBER: US 08/027,146

PRIOR APPLICATION NUMBER: US 08/027,146

PRIOR APPLICATION NUMBER: US 08/073,205

PRIOR APPLICATION NUMBER: US 08/073,396

PRIOR APPLICATION NUMBER: US 08/103,396

PRIOR APPLICATION NUMBER: US 08/159,184

PRIOR APPLICATION NUMBER: US 08/159,184

PRIOR APPLICATION NUMBER: US 08/159,399

PRIOR APPLICATION NUMBER: US 08/159,399

PRIOR APPLICATION NUMBER: US 08/159,339

PRIOR APPLICATION NUMBER: US 08/159,339

PRIOR APPLICATION NUMBER: US 08/205,713

PRIOR APPLICATION NUMBER: US 08/205,713

PRIOR APPLICATION NUMBER: US 08/205,713

PRIOR APPLICATION NUMBER: US 08/347,610

PRIOR FILING DATE: 1994-03-04

PRIOR APPLICATION NUMBER: US 08/347,610
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; LENGTH: 7
; TYPE: PRT
; TYPE: PRT
; ORCANISM: Bovine
US-10-494-781-6
RESULT 15

US-11-045-024-3575

; Sequence 3575, Application US/11045024

; Publication No. US20050271676A1

; GENERAL INFORMATION:
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US-11-045-024-3529
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SEQ ID NO 3529
LENGTH: 9
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Best Local
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Best Local Similarity
Matches 4; Conserv
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hes 5; Conserv
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Denise Marie
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100.0%; Pred. No. 5.8
tive 0; Mismatches
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    Mismatches

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Pred. No. 5.8e+04;
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5.8e+04;
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PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-06-04
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR FILING DATE: 1993-11-29
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APPLICANT: Epimmune Inc.

TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007

CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
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APPLICANT:
APPLICANT:
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Livingston, Brian
APPLICANT: Chesnut, Robert
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SOFTWARE: FastSEQ for Windows Version 4.0
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TYPE: PRT
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CURRENT FILING DATE: 2005-01-28
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Celis, Esteban
Kubo, Ralph
Grey, Howard M.
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Baker, Denise Marie
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Pred. No. 5.
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PRIOR FILING DATE: 1993-11-29

PRIOR APPLICATION NUMBER: US 08/205,713

PRIOR FILING DATE: 1994-03-04

PRIOR APPLICATION NUMBER: US 08/347,610

PRIOR FILING DATE: 1994-12-01

NUMBER OF SEQ ID NOS: 14528

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 10581
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US-11-045-024-10581
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; TYPE: PRT
; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
US-11-045-024-3576
                                          US-11-045-024-10581
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PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3576
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Publication No. US20050271676A1
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Query Match
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PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,339
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PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-06-04
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CURRENT FILING DATE: 2005-01-28
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TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
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                                                             ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
                                                                                                 ENGTH: 9
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OR APPLICATION NUMBER: US 08/103,396
OR FILING DATE: 1993-08-06
OR APPLICATION NUMBER: US 08/159,184
OR FILING DATE: 1993-11-29
OR APPLICATION NUMBER: US 08/159,339
OR FILING DATE: 1993-11-29
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Similarity 62.5%;
5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Robert
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  34.4%; Score 22;
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Pred. No. 5.8e+04;
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  7;
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Length 9;
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APPLICANT:

APPLICANT:

APPLICANT:

Livingston, Brian Chesnut, Robert Baker, Denise Marie Celis, Esteban

APPLICANT:

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Query Match
Best Local Similarity
Thes 5; Conserve
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                                                                                                                      RESULT 19
US-11-045-024-10720
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US-11-045-024-10652
GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
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SEQ ID NO 10652
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Publication No. US20050271676A1
                                                                                Sequence 10720, Application US/11045024 Publication No. US20050271676A1
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TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007
CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 1999-10-05
PRIOR PRIOR APPLICATION NUMBER: US 08/071,146
PRIOR FILING DATE: 1993-03-05
PRIOR FILING DATE: 1993-08-04
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PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/205,713
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PRIOR APPLICATION NUMBER: US 08/347,610
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Celis, Esteban
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1; Mismatches 3,
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Pred. No. 5.8e+04;
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Sequence 12381, Application US/11045024
Publication No. US20050271676A1
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              TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions FILE REFERENCE: 2060.0040007
CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR PILING DATE: 1999-10-05
PRIOR PILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/03,205
PRIOR APPLICATION NUMBER: US 08/03,305
PRIOR APPLICATION NUMBER: US 08/03,305
PRIOR APPLICATION NUMBER: US 08/03,305
PRIOR APPLICATION NUMBER: US 08/03,306
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-08-06
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SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR FILING DATE: 1993-03-05
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TYPE: PRT
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PRIOR APPLICATION NUMBER: US 08/159,339
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PRIOR FILING DATE: 1993-06-04
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILING DATE: 1993-08-06
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APPLICATION NUMBER: US 08/159,184
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5; Conserv
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Pred. No. 5.
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US-11-045-024-12445
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                               Matches
                                                Query Match
Best Local
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PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR
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PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
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CURRENT FILING DATE: 2005-01-28
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                                                                                                      TYPE: PRT
ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
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FILING DATE: 1993-08-06
APPLICATION NUMBER: US 08/159,184
FILING DATE: 1993-11-29
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LALVRELK 13
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Grey, Howard M.
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Chesnut, Robert
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                                               Score 22; DB 7;
Pred. No. 5.8e+04
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Pred. No. 5.8e+04;
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RESULT 23
US-11-045-024-13980
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PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
                                                                                                                                                                                     APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Livingston, Brian
APPLICANT: Chesnut, Robert
APPLICANT: Baker, Denise Mari
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Best Local Similarity 62.5%;
Matches 5; Conservative
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              APPLICANT: Kubo, Ralph
APPLICANT: Grey, Howard M.
APPLICANT: Epimmune Inc.
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007
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PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-06-04
PRIOR FILING DATE: 1993-06-04
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CURRENT APPLICATION NUMBER: US/11/045,024
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                                                                                                                                                             Livingston, Brian
Chesnut, Robert
Baker, Denise Marie
Celis, Esteban
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Baker, Denise Marie
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; Pred. No. 5.8e
1; Mismatches
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5.8e+04;
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APPLICANT: Epimmune Inc.
TITIE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITIE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007
CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/03,205
PRIOR FILING DATE: 1993-08-04
PRIOR PILING DATE: 1993-08-06
PRIOR PILING DATE: 1993-08-06
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,184
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US-11-045-024-3581
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILLING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILLING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILLING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3581
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PRIOR FILLING DATE: 1993-06-04
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILLING DATE: 1993-08-06
PRIOR FILLING DATE: 1993-08-06
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR FILLING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILLING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILLING DATE: 1994-03-04
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR FILING DATE: 1993-03-05
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TYPE: PRT
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Local Similarity 55.6%;
les 5; Conservation
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Celis, Esteban
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                               for Windows Version 4.0
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Pred. No. 5.8e+04;
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CURRENT ETLING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR FILING DATE: 1993-06-04
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILING DATE: 1993-10-06
PRIOR FILING DATE: 1993-11-29
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
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SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 3582
LENGTH: 10
TYPE: PRT
ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
US-11-045-024-3582
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TYPE: PRT
; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
US-11-045-024-3581
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US-11-045-024-3582
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TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.00440007
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3 LELLEELK 10
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Southwood, Scott
Livingston, Brian
Cheemut, Robert
Baker, Denise Marie
Celis, Esteban
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progesterone recep
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hypothetical prote

galactose-1-phosph T-cell receptor ga acidic fibroblast

hypothetical

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hypothetical prote insertion element Ig heavy chain DJ Ig heavy chain CRD glycophorin B (mis Ig heavy chain V r hemocyanin chain S Sf11-dlycoprotein

nuclear antigen EB T cell receptor al

protein QA100044 stylar glycoprotei Ig H chain V-D-J r D-galactose-bindin protein kinase C i

phenylalanine-tRNA hypothetical 2.1K

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R;Nyitray, L.; Mocz, G.; Szilagyi, L.; Balint, M.; Lu, R.(J. Biol. Chem. 258, 13213-13220, 1983
A;Title: The proteolytic substructure of light meromyosin. A;Reference number: A92403; MUID:84032553; PMID:6355107
A;Accession: A20923
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         globin - polychaete (Eudistylia v
N;Alternate names: chlorocruorin
C;Species: Eudistylia vancouveri
                                                                                                                                                                                                                                                       light meromyosin - chicken (fragment)
C;Species: Gallus gallus (chicken)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 31-Dec-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Qabar, A.N.; Stern, M.S.; Walz, D.A.; Chiu, J.T.; Timkovich, R.; Wall, J.S.; Kapp, J. Mol. Biol. 222, 1109-1129, 1991
A;Title: Hierarchy of globin complexes. The quaternary structure of the extracellula: A;Reference number: S19532; MUID:92106333; PMID:1762147
A;Accession: S19615
A;Cross-references: UNIPROT:Q8JG72; UNIPARC:UPI00001775EC C;Superfamily: myosin motor domain homology C;Keywords: ATP; muscle
                                                                   A; Molecule type: prote
A; Residues: 1-18 <NYI>
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A;Experimental source: plume
C;Complex: dodecamers, each consisting of a trimer of tetramers of globin chains; dodecamer: neme; homotetramer; oxygen carrier
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A;Residues: 1-20 <QAB>
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5.1e+02;
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A;Gene: rplJ
C;Superfamil
C;Keywords: |
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S35970
                            hydroxypyruvate reductase (EC 1.1.1.81) - Methylobacterium extorquens (fragment) C;Species: Methylobacterium extorquens C;Date: 01-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 08-Oct-1999 C;Accession: A44921
                                                                                                                                                                                                                                                                                                                                                                                               submitted to the EMBL Data Library, A;Reference number: S35970 A;Accession: S35978
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C;Accession:
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C;Species: Proteus vulgaris
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A; Residues: 1-21 < ZHY>
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A; Residues: 1-20 <ZHY>
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                                                                                                      RESULT 6
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A;Cross-references: UNIPROT:Q7M532; UNIPARC:UPI000017AEA3
R;Engemann, S.; Noelle, R.; Herfurth, E.; Briesemeister, U.; Grelle
Eur. J. Blochem. 234, 24-31, 1995
A;Title: Carcography of ribosomal proteins of the 30S subunit from
A;Reference number: S63964; MUID:96098717; PMID:8529646
A;Accession: S63967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 7
PA0015
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A;Molecule type: DNA
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-20 <CHI>
A;Residues: 1-20 <CHI>
A;Cross-references: UNIPARC:UPI000016F931; GB:M81443; NID:g150010; PIDN:AAA25378.1;
A;Note: sequence extracted from NCBI backbone (NCBIN:75202, NCBIP:75203)
C;Keywords: oxidoreductase
                                                                                                                                                                                                                                                                                                                               R;Engemann, S.; Herfurth, E.; Briesemeister, U.; Grelle, G.; Wittmann-Liebold, submitted to the Point Sequence Database, November 1994
A;Description: Cartography of ribosomal proteins of the 30S subunit from the harmaneric number; S49023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Experimental source: seed
C;Keywords: pyroglutamic acid; seed; storage protein
F;1/Modified site: pyrrolidone carboxylic acid (Gln)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 1-14 <KAM>
A;Cross-references: UNIPARC;UPI000017B010
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A; Reference number:
A; Contents: AM1
                                                                                                   A; Molecule type: protein A; Residues: 1-18 < ENW>
                                                                                                                                                                                                                                                                                                                                                                                                                                ribosomal protein HS25 [validated] - Haloarcula marismortui (fragment)
C;Species: Haloarcula marismortui
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
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A;Accession: PA0015
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A;Residues: 1-18 <ENG>
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Best Local S
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Query Match
Best Local Similarity
                                                        Cross-references: UNIPARC:UPI000017AEA3
Keywords: blocked amino end; protein biosynthesis; ribosome
                                                                                                                                                                                                                                                                                                                                                                                                                   Accession: S49026; S63967
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32.8%;
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Score
Pred.
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Pred. No. 1.3e+03;
3; Mismatches 4
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Pred. No. 1.2e+03;
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2.5e+03;
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C; Keywor
                                                          R; Uezono, Y.; Tsumori, H.; Mukasa, submitted to JIPID, October 1993
                                                                                               dextransucrase (EC 2.4.1.5) Streptococcus bovis (fragment) C;Species: Streptococcus bovis C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_chaC;Accession: PU0034
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                   A; Reference number:
                                      A; Description: Purification and properties
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A; Residues: 1-20 < MAR>
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                       PU0034
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Biochem. J. 300, 567-575, 1994
A;Title: Novel inducible antibacterial peptides from a hemipteran insect, the
A;Reference number: S44463; MUID:94271176; PMID:8002963
A;Accession: S44465
                                                                                                                                                                                                                                                                                                                                              pyrrhocoricin - Pyrrhocoris apterus
C;Species: Pyrrhocoris apterus
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Martin, B.; Sharples, G.J.; Humbert, O.; Lloyd, R.G.; Claverys, J.P. Mol. Microbiol. 19, 1035-1045, 1996
A;Title: The mmsA locus of Streptococcus pneumoniae encodes a RecG-like protein A;Reference number: S71015; MUID:96249697; PMID:8830261
A;Accession: S71017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein - Streptococcus pneumoniae (fragment)
C;Species: Streptococcus pneumoniae
C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_c
C;Accession: S71017
                                                                                                A;Description: antibacterial protein A;Note: active against Gram-negative
                                                                                                                                                                  A;Cross-references: UNIPROT:P37362; UNIPARC:UPI0000132E13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary; nucleic acid sequence
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                                                                               antibacterial; hemolymph;
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66.7%;
                 Score 21;
Pred. No. :
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Pred. No. 2.7e+03;
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                                     Length 20;
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glucosyltransferase

synthesizing 1,6-alph

#text_change

09-Jul-2004

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RESULT 14
A35550
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c;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 09-Jul-2004
C;Accession: I55226
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A; Residues: 1-11
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A;Residues: 1-17 <BLO9
A;Cross-references: UNIPROT:P14460; UNIPARC:UPI000012A764
A;Cross-references: UNIPROT:P14460; UNIPARC:UPI000012A764
A;Blombaeck, B.; Blombaeck, M.; Groendahl, N.J.
Rcta Chem. Scand. 19, 1789-1791, 1965
A;Tille: Studies on fibrinopeptides from mammals.
A;Reference number: A03118
                                                                                                                                                                                                                                                                                                                                                                                   unpublished results, cited A;Reference number: A29501 A;Accession: E29501
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R;Blombaeck, B.; Blombaeck,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Sus scrofa domestica (domestic pig)
C;Date: 21-Nov-1987 #sequence_revision 21-Nov-1987 #text_change 09-Jul-2004
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E29501
fibrinopeptide A - pig
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lt fast myosin heavy chain from mouse skeletal muscle A;Reference number: I55226; MUID:84061805; PMID:61963: A;Accession: I55226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Weydert, A.; Daubas, P.; Caravatti, M.; Minty, A.; Bugaisky, G.; Cohen, A.;
J. Biol. Chem. 258, 13867-13874, 1983
A;Title: Sequential accumulation of mRNAs encoding different myosin heavy cha:
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A; Residues: 1-17 < RES>
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Best Local
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3; Conserv
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Similarity 36.4%;
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57.1%;
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Pred. No. 2.3e+03;
                                                                                                                                            Score 20; DB 2;
Pred. No. 3.5e+03;
1; Mismatches
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Pred. No.
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R;Shigeoka, S.; Nakano, Y.
Arch. Biochem. Biophys. 288, 22-28, 1991
A;Title: Characterization and molecular properties of 2-oxoglutarate decarboxylase
A;Reference number: A61150; MUID:91378443; PMID:1910306
A;Accession: A61150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Superfamil
C;Keywords:
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B53875
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A;Molecule type: protein
A;Residues: 1-17 <SHI>
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R;Shi, D.L.; Savona, C.; Gagnon, J.; Cochet, C.; Chambaz, E.M.; Feige, J.J.
J. Biol. Chem. 265, 2881-2887, 1990
A;Title: Transforming growth factor-beta stimulates the expression of alpha-2-macroglobu A;Reference number: A35550; MUID:90153919; PMID:1689294
A;Accession: A35550
                                                                                                                                                                                                                                                                                                                                                                                  RESULT
A61150
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A;Experimental source: Brockmann body, principal islet
A;Note: sequence extracted from NCBI backbone (NCBIP:120600)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            J. Protein Chem. 11, 489-494, 1992
A;Title: The principal islet of the Coho salmon (Oncorhyncus kisutch)
A;Reference number: A53875; MUID:93080727; PMID:1449598
A;Accession: B53875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 adrenocortical cell major secreted protein - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Date: 21-Sep-1990 #sequence_revision 21-Sep-1990 #text_change
                                                                                                                                                                                                                                                                                                         2-oxoglutarate decarboxylase (EC 4.1.1.71) - Euglena gracilis (fragment)
C;Species: Euglena gracilis
C;Species: 10-Mar-1994 #text_change 09-Jul-2004
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C;Species: Oncorhynchus kisutch (coho salmon)
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                                                                                                              A;Cross-references: UNIPROT:Q7M266; UNIPARC:UPI000017AED8
                                                                                                                                     A;Molecule type: protein
A;Residues: 1-20 <SHI>
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                                                                                            ;Keywords:
                                                                                                                                                                                                                                                                                           Accession: A61150
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    Matches
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Best Local
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Pred. No. 3.5e+03;
1; Mismatches 5
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Pred. No. 4.1e+03;
                                              Length 20;
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YLALVRELK 13

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glucocerebrosidase - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #sequence_revisio
C;Accession: I52980; I65971
                                                                               RESULT 19
I52980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   glutathione transferase (EC 2.5.1.18) class alpha 5 - pig (fragment)
N;Alternate names: glutathione S-transferase class alpha 5
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 19-Mar-1998 #sequence_revision 17-Apr-1998 #text_change 07-May-1999
C;Accession: S71867
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI00001754D3
C;Comment: At least five speciés-independent classes of cytosolic glutathion s mitochondrial form are known.
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Biochem. J. 317
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A;Title: A new Tetrahymena actin-binding protein is localized A;Title: A new Tetrahymena actin-binding protein is localized A;Reference number: PD0015; MUID:98207047; PMID:9538250
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A;Residues: 1-21 <WAT>
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Date: 13-Jun-1998 #sequence_revision 10-Jul-1998 #text_change 09-Jul-2004
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Best Local
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ilarity 75.0%;
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3; Mismatches
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                  #text_change 05-Nov-1999
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A;Cross-references: UNIPARC:UPI00001427C8
A;Experimental source: style, strain Mill
R;Anderson, M.A.; Cornish, E.C.; Mau, S.L.; William
J.P.; Crawford, R.J.; Clarke, A.E.
Nature 321, 38-44, 1986
A;Title: Cloning of cDNA for a stylar glycoprotein
A;Reference number: A28819
                                                                                                                                                                                                                                                 C;Species: Lycopersicon peruvianum (Peruvian tomato)
C;Date: 23-Nov-1991 #sequence_revision 23-Nov-1991 #
C;Accession: PQ0193; A28819
R;Mau, S.L.; Williams, E.G.; Atkinson, A.; Anderson,
Planta 169, 184-191, 1986
                                                                                                                                                 A; Molecule type: protein A; Residues: 1-15 < MAU>
                                                                                                                                                                                                            A; Title: Style proteins of A; Reference number: PQ0192
                                                                                                                                                                                                                                                                                                                                                           stylar glycoprotein 3 - Peruvian tomato
N;Alternate names: S3-glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEBS Lett. 317, 53-56, 1993
A;Title: Association of three small GTP-binding proteins with A;Reference number: S29485; MUID:93154521; PMID:8428634
A;Accession: S29486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTP-binding protein o-rab3 . C; Species: Discopyge ommata
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DNA 7, 107-116,
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R;Volknandt, W.; P
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A; Residues: 1-11 <RE2>
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A; Residues: 1-11 < RES>
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42.9%;
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Pred. No. 4.4e+03;
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Pred. No. 3.5e+03
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RESULT 24
PH1778
T cell receptor alpha chain V region (clone 1PBL V alpha 24-5) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1778
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A;Cross-references: UNIPARC:UPI00001427C8
C;Keywords: glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: S10809
R;Toker, A.; Ellis, C.A.; Sellers, L.A.; Aitken, A.
Eur. J. Biochem. 191, 421-429, 1990
A;Title: Protein kinase C inhibitor proteins. Purification from sheep brain and sequence A;Reference number: S10804; MUID:90345949; PMID:2143472
A;Accession: S10809
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein kinase C inhibitor KCIP-1 isoform f - sheep (fragment)
N;Alternate names: probable lipocortin
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 19-Mar-1997 #sequence_revision 13-Sep-1998 #text_change 13-Sep-1998
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allelic exclusion and antigen-specific repertoire.
A,Reference number: PH0746; MUID:92078846; PMID:1836010
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0784
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A; Residues: 1-15 < CAS>
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J. Exp. Med. 174, 1371-1383, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T-cell receptor alpha chain (F1) - mouse (fragment)
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Local Similarity 66.7%;
ss 4; Conservation
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Pred. No. 4.7e+03;
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Pred. No. 4.7e+03;
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Pred. No. 5e+03;
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Search completed: January 26, Job time: 9.5 secs

2006, 08:05:00

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RESULT 25
PH1781
T cell receptor alpha chain V region (clone 2DN V alpha 24-2) -
C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change
C;Accession: PH1781
                                                                                                                                             A;Molecule type: mRNA
A;Residues: 1-16 <POR>
A;Cross-references: UNIPARC:UPI000017C374
                                                                                                                                                                                                A;Reference number: PH1754; MUID:93301585; PMID:8391057
A;Accession: PH1781
A;Status: preliminary
                                                                                                                                                                                                                                                                         R;Porcelli, S.; Yockey, C.E.; Brenner, M.B.; Balk, S.P. J. Exp. Med. 178, 1-16, 1993
A;Title: Analysis of T cell antigen receptor (TCR) expression
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J. Exp. Med. 178, 1-16, 1993
A;Title: Analysis of T cell antigen receptor (TCR) expression by human peripheral blood
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A;Cross-references: UNIPARC:UPI000017C36C
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A;Residues: 1-16 <POR>
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Pred. No. 5e+03;
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Q4KN88
WUSSP PRELIMINARY;
Q4KN88;
13-SEP-2005 (TremBLrel. 31,
13-SEP-2005 (TremBLrel. 31,
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PROTEIN SEQUENCE.

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PRODE STATE OF THE PROPERTY OF THE PROTEIN SEQUENCE.

MUCH Lendon G. Witt H.T., Herrmann R.G.;

"Genes encoding eleven subunits of photosystem I from the thermophicyanobacterium Synechococcus sp.";

Gene 127:71-78(1993).

GO; GO:0010979; P:photosystem I; IEA.

GO; GO:0011979; P:photosynthesis; IEA.

InterPro; IPR010010; PsaM.

Pfam; PF07465; PsaM; 1.

SEQUENCE 20 AA; 2102 MW; DF585D9ED9D7C512 CRC64;
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Bacteria; Cyanobacteria;
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01-MAY-2000 (TrEMBLrel. 13, Last seque
01-MAR-2004 (TrEMBLrel. 26, Last annot
Photosystem I subunit XII (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUN-2002 (TrEMBLrel. Flagellin (Fragment). Proteus mirabilis.
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MEDLINE=91372967; PubMed=1680106;
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Enterobacteriaceae; Proteus.
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Query Match
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01-DEC-1992 (Rel. 2
01-DEC-1992 (Rel. 2
05-JUL-2004 (Rel. 4
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammaalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Guidry P.A., Stroynowski I.;
"The Murine Family of Gut-Rest:
Alternatively Spliced Isoforms
Blastocyst MHC.";
                                                                                                                        This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                         J. Bacteriol. 173:6844-6848(1991).

-!- CATALYTIC ACTIVITY: Isocitrate = succinate + glyoxyl-
-!- PATHWAY: Glyoxylate bypass; first step.
-!- SUBUNIT: Homotetramer (By similarity).
-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the isocitrate lyase family.
                                                                                                                                                                                                                                                                                                                                                      PROTEIN SEQUENCE.
MEDLINE=92041568; PubMed=1938889;
Hoyt J.C., Johnson K.E., Reeves H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NON TER
InterPro, IPR000918; Isocit lyase ph. PROSITE; PS00161; ISOCITRATE LYASE; Pr Direct protein sequencing; GTyoxylate Tricarboxylic acid cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Acinetobacter calcoaceticus.
Acinetobacter calcoaceticus.
Racteria; Proteobacteria; Gammaproteobacteria;
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                                                                      PIR; A41338; A41338.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria; Proteobacteria; Gan
Moraxellaceae; Acinetobacter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isocitrate lyase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (MAR-2005) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=SEB;
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                                                                                                                                                                                                                                                                                                                 socitrate lyase."
                                                                                                                                                                                                                                                                                                                                  "Purification and characterization
                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=471;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Fragment).
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. 24, Last sequence update)
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(EC 4.1.3.1) (Isocitrase) (Isocitratase)
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                                      PARTIAL
                    bypass;
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83AE726B1F2F96E3 CRC64;

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Score 25; Pred. No.

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RESULT 5
O7M3Q5_EUDVA
O7M3Q5_EUDVA PRELIMINARY;
AC Q7M3Q5;
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Globin (Fragment).
OS Euddstylia vancouverii.
OC Eukaryota; Metazoa; Annelid.
OC Eukaryota; Metazoa; Annelid.
OC Sabellida; Sabellidae; Eudi.
OX NCBL_TAXID=6364;
RN [1]
RR PROTEIN SEQUENCE.
RR MEDLINE=92106333; PubMed=17,
RA Qabar A.N., Sterm M.S., Wal.
RA Kapp O.H., Vinogradov S.N.;
RT "Hierarchy of globin comple.
RT extracellular chlorocruorin
RL J. Mol. Biol. 222:1109-1129
DR PIR; S19615; S19615.
FT NON_TER 20
SEQUENCE 20 AA; 2286 MW;
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Q76UT2_9CAUD
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Best Local S
Matches 5
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SEQUENCE
                                                                                                                                                           MEDLINE-96064424; PubMed-8526525;
Boyce J.D., Davideon B.E., Hillier A.J.;
"Spontaneous deletion mutants of the Lactococcus lactis temperate bacteriophage BK5-T and localization of the BK5-T attP site.";
Appl. Environ. Microbiol. 61:4105-4109(1995).
EMBL; E80308; AAB35788.2; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                            05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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01-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                       Lactococcus phage BK5-T.
Viruses; deDNA viruses,
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Q76UT2;
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Kapp O.H., Vinogradov S.N.,
                                                                                                                                                                                                                                                                                                                               Integrase (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hierarchy of globin complexes. The quaternary structure extracellular chlorocruorin of Eudistylia vancouverii."; Mol. Biol. 222:1109-1129(1991).
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Sabellidae; Eudistylia.
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M.S., Walz D.A.,
                                                                                                                                    1831 MW;
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41.7%;
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Pred. No. 3.6e
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3.6e+03;
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Best Local (
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01-MAR-2001
01-JUN-2003
                                   NUCLEOTIDE SEQUENCE.
STRAIN=Z-G T95-873;
MEDLINE=99175282; PubMed=10074533;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WEDLINE=97448947; PubMed=9303271; DOI=10.1016/S0145-305X(97)00012-8; Wittwer D., Weise C., Goetz P., Wisemer A.; "LPS (lippopolysaccharide)-activated immune responses in a hemocyte cell line from Estigmene acraea (Lepidoptera)."; Dev. Comp. Immunol. 21:323-336(1997)

Dev. Comp. Immunol. 21:323-336(1997)

-i- FUNCTION: Lysozymes have primarily a bacteriolytic function; those in tissues and body fluids are associated with the monocyte-macrophage system and enhance the activity of immunoagents (By
                                                                                                        Human adenovirus 4 (HAdV-4)
Viruses; dsDNA viruses, no
                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR001916; Glyco_hydro_22.
prOSITE; pS00128; LACTALBUMIN_LYSOZYME; PARTIAL.
Antimicrobial; Bacteriolytic enzyme; Direct protein sequencing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-OCT-2001
16-OCT-2001
           Crawford-Miksza L.K., Nang R.N., "Strain variation in adenovirus
                                                                                         NCBI_TaxID=28280;
                                                                                                                                              P23 protease
                                                                                                                                                                                                      Q9DSS9;
                                                                                                                                                                                                                   Q9DSS9_ADE04 PRELIMINARY;
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13-SEP-2005 (Rel. 48, Last annotation update)
Lysozyme (EC 3.2.1.17) (1,4-beta-N-acetylmuramidase)
Estigmene acrea (Salt marsh moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pt
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European
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SIMILARITY: Belongs to the glycosyl hydrolase 22 family.
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                                                                                                        Adenoviridae; Mastadenovirus.
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RESULT 10
Q4XBH7 PLA
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Best Local S
Matches 5
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Matches 6
BH7_PLACH PRELIMINARY;
Q4XBH7_PLACH PRELIMINARY;
Q4XBH7_13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Crawford-Miksza L.K.;
Submitted (DEC-2000) to the
SMBL; AF065064; AAD03661.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
STRAIN=Z-G T95-873;
                                                                                                                                                                               INIT MET
                                                                                                                                                                                                                                                                                                                                      Zhyvoloup A.N.;
Submitted (AUG-1993) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the ribosomal protein L10P family.
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P43448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                               This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriaceae; Citrobacter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          50S ribosomal protein
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                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                 PROSITE; PS01109; RIBOSOMAL RIBOSOMAL RIBOSOMAL RIBOSOMAL
                                                                                                                                                                                                                               HAMAP; MF_00362; -; 1
                                                                                                                                                                                                                                             EMBL; X74448; CAA52459.1; -; PIR; S35970; S35970.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Citrobacter freundii.
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13-SEP-2005
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6; Conserv
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                                                                                     DKQAIVAEVSEV 17
                                                                                                         DKQNYLALVREL 12
                                                                                                                                                                                                              IPR002363; Ribosomal_L10eub.
S01109; RIBOSOMAL_L10; PARTI
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                                                                                                                                                                     19 AA;
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(Rel. 32,
(Rel. 48,
  (TrEMBLrel. 31, (TrEMBLrel. 31,
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8 AA;
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1917 MW;
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45.5%;
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Last annotation updat
L10 (Fragment).
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            Created)
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-; Genomic_DNA.
                                                                                                                            Score 24; DB Pred. No. 4.66
2; Mismatches
                                                                                                                                                                                        protein.
By simil
                                                                                                                                                                                                                                                       Genomic_DNA.
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Pred. No.
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                                                                                                                                                                                        similarity.
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                                                                                                                                                                                                              PARTIAL.
   sequence update
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RESULT 11
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Matches
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01-OCT-1996
13-SEP-2005
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                                                                                                                              INIT MET
NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zhyvoloup A.N.;
Zhyvoloup A.N.;
Submitted (AUG-1993) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the ribosomal protein L10P family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RL10_PROVU
P51411;
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Eukaryota; Alveolata;
NCBI_TaxID=5825;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                 HAMAP; MF_00362; -; 1.
InterPro; IPR002363; Ribosomal_L10eub.
PROSITE; PS01109; RIBOSOMAL_L10; PARTIAL.
                                                                                                                                                                                                                                                                                                                                                                EMBL; X74446; CAA52457.1; -; PIR; S35978; S35978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its tuse as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteria; Proteobacteria; Ga
Enterobacteriaceae; Proteus.
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                                                                                                                                                                                                                                     Ribonucleoprotein; Ribosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=585;
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6,
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(Rel. 34, Last sequence update)
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Conservative
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0 AA;
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2083 MW;
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By simil
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Pred. No. 4.6e+0
                            Score 24; DB 1;
Pred. No. 4.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                   Genomic_DNA.
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                                                                                                                                  76520A02CD7851DB
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   Mismatches
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                                                               Length
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DKQNYLALVREL

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ID 13-SEP-2
ID 13
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NON TER
SEQUENCE
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MEDLINE=22306558; PubMed=12420160; DOI=10.1007/800203-002-0458-7;

RYChlik I., Martin G., Methner U., Lovell M., Cardova L., Sebkova A.,

Sevcik M., Damborsky J., Barrow P.A.;

"Identification of Salmonella enterica serovar Typhimurium genes

associated with growth suppression in serionary-phase nutrient broth

cultures and in the chicken intestine.";

Arch. Microbiol. 178:411-420(2002).

EMBL; AF268390; AAF73476.1; -; Genomic_DNA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0015288; F:porin activity; IEA.

GO; GO:0015288; F:porin activity; IEA.
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-I- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is northing to the sequence shown here is derived from an entry which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Putative glycoporin (Fragment). Salmonella typhimurium.
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01-OCT-2000 (TrEMBLrel. 15,
01-JUN-2003 (TrEMBLrel. 24,
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MEDLINE=93184488; PubMed=1292782;
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01-MAR-2002 (TrEMBLrel. 20
01-MAR-2002 (TrEMBLrel. 20
01-UN-2003 (TrEMBLrel. 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NADH dehydrogenase subunit 1 (Fragment) Discorsopagurus schmitti. Mitochondrion.
                                                           Submitted (GEP-1997) to the EMBL/GenBank/DDBJ da EMBL; AF022930; AAD01753.1; -; Genomic_DNA.
GO; GO:0008233; F:peptidase activity; IEA.
GO; GO:0006508; F:proteolysis and peptidolysis; InterPro; IPR005082; P:proteolysis and peptidolysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q9ZXI2 BPAR1 PRELIMINARY;
Q9ZXI2;
Q1-MAY-1999 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Anomura; Paguroidea;
Paguridae; Discorsopagurus.
                                                                                                                                                                                                                                                                                                                                   Ompc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Viruses; dsDNA v
T4-like viruses.
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       NON
                          Protease.
                                               Pfam; PF03420; Peptidase_U9;
                                                                                                                                                                                          Syu W.J., Ding H.C., Chang Y.C.;
                                                                                                                                                                                                                                        STRAIN-AR1
                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                      "Characterization of a 0157:H7 and disclosure
                                                                                                                                                                                                                                                                                                                                                         O157:H7
                                                                                                                                                                                                                                                                                                                                                                                                    Syu W.;
                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=98433034; PubMed=9758911; DOI=10.1159/000025351; Yu S., Ding H., Seah J., Wu K., Chang Y., Chang K.S., Ta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-AR1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=g21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
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1878 MW;
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66.7%;
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No. 6.2e+03;
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M.,
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in host outer membrane protein
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RESULT 18
Q77WL5_ADE04
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AC CRAWFORD_ADE04
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Q9W9C1_ADI
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Best Local
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01-MAR-2001
01-JUN-2003
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Q9W9C1_.
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MEDLINE=99175282; PubMed=10074533;
Crawford-Miksza L.K., Nang R.N., Schnurr [
"Strain variation in adenovirus serotypes"
                                                                                                                                           Human adenovirus 4 (HAd
Viruses; dsDNA viruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Crawford-Miksza I.K.;
Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065063; AAD03658.3; -; Genomic_DNA.
GO; GO:00004197; F:cysteine-type endopeptidase activity; IEA.
GO; GO:0008233; F:peptidase activity; IEA.
GO; GO:0008233; F:proteolysis and peptidolysis; IEA.
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Viruses; dsDNA viruses, no
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05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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                    MEDLINE=99175282; PubMed=10074533; Crawford-Mikeza L.K., Nang R.N., S
                                                            NUCLEOTIDE SEQUENCE.
STRAIN=RI-67;
                                                                                                                                                                                                                                                                                    Q9W9C1;
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J. Clin. Microbiol. 37
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                                                                                                                    NCBI_TaxID=28280;
                                                                                                                                                                                                     P23 protease
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ProDom; PD003705; Peptidase_C5; 1.
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    variation
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                                                                                                                                                                                                     (Fragment)
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.K., Nang R.N., Schnurr D.P.;
in adenovirus serotypes 4 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1929
                                                                                                                                                                  (HAdV-4)
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Q6LBF5 HUMAN
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Crawford-Miksza L.K.;

Crawford-Miksza L.K.;

Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.

Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.

GO; GO:0004197; F:cysteine-type endopeptidase activity;

GO; GO:0006203; F:peptidase activity; IEA.

GO; GO:0006508; P:proteolysis and peptidolysis; IEA.

InterPro; IPR000855; Peptidase C5.

ProDom; PD003705; Peptidase C5.
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05-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (JUL-1997) to the EMBL/GenBank/DDBJ databases. EMBL; X77531; CAA54669.1; -; Genomic DNA.
GO; GO:0030195; P:negative regulation of blood coagulation of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Euarchonto
                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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J. Clin. Microbiol. 37
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NUCLEOTIDE SEQUENCE Hall N., Karras M.,
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                                      Hypothetical protein (Fragment)
ORFNames=PC400809.00.0;
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Q4XIZ2;
Eukaryota; Alveolata;
NCBI_TaxID=5825;
                                                                                   13-SEP-2005
13-SEP-2005
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Cook C.E., Smith M.L., Telford M.J.,
"Hox genes and the phylogeny of the
Curr. Biol. 11:759-763 (2001).
                             Plasmodium chabaudi
                                                                     13-SEP-2005
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01-MAR-2001 (TrEMBLrel. 16, Last sequence up
01-DEC-2001 (TrEMBLrel. 19, Last annotation
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01-MAR-2001
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Pleurostigmophora; Lithobiomorpha; Lithobiidae; Litho
NCBI_TaxID=7552;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Scutigerella.
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Eukaryota; Metazoa; Arthropoda;
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Scutigerella immaculata (gar
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EMBL; AF362097; AAK51952.1; -; Genomic_DNA.
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Engrailed (Fragment).
Lithobius forficatus.
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                                                                                                                                                                                                                                                                                                                                                                             PubMed=11378385; DOI=10.1016/S0960-9822(01)00222-6; M.L., Telford M.J., Bastianello A., Akam M.;
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Pred. No. 7e+03;
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Pred. No. 7e+03;
D; Mismatches
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RESULT

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OBWGC5 9EUCA
OBWGC5-9EUCA PRELIMINARY; PRT; 20 AA.
ID QBWGC5;
DT 01-MAR-2002 (TrEMBLrel. 20, Created)
DT 01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE NADH dehydrogenase subunit 1 (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE.
Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Hall N., Karras M., Florens L., Janssen C.S., Pain A., Christophides G.K.,
Berriman M., Florens L., Janssen C.S., Pain A., Churcher C.,
James K., Rutherford K., Harris B., Harris D., Churcher C.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.C.,
Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.,
"A comprehensive survey of the Plasmodium life cycle by genomic,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plasmodium chabaudi.
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
NCBI_TaxID=5825;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transcriptomic, and proteomic analyses.";
Science 307:82-86(2005).
-I- CAUTION: The sequence shown here is derived
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PLACH
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EMBL; AF436053; AAL31625.l; -; Ge
GO; GO:0005739; C:mitochondrion;
                                                                                                                                                                            MEDLINE=21884466; PubMed=11886621; Morrison C.L., Harvey A.W., Lavery Cunningham C.W.;
                                                                                                                                                                                                                                               Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Anomura; Hippoidea;
Albuneidae; Blepharipoda.
                                                                                                                                                                                                                                   NCBI_TaxID=177216;
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                                                                                                                                             269:345-350(2002).
                                                                                                                                                                                         DOI=10.1098/rspb.2001.1886;
S., Tieu K., Huang Y.,
                                                                                                                                                                     confirm the
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Search completed: January 26, 2006, 08:03:51 Job time: 41.6638 secs

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   DMAQNYKYRQQFIQSVLNNGATRQ
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Copyright (c) 1993 - 2006 Compugen
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ABA01731

ABA01731

ABG96178

ADM115661

ABY01496614

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ABR31386

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Adm53387 Beta-cate
Adm95021 Wild-type
Adm95021 Wild-type
Adj78231 Peptide S
Adr31521 Beta-cate
Aea01731 Human bet
Abg96178 Cysteine-
Aea01731 Gliadin r
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10-APR-2001;
25-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcripts, and as therapeutic agents. The proteins and peptides are useful as therapeutic prognostic and diagnostic reagents for cancer. The present sequence is a human leukocyte antigen (HLA) peptide, used in an example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to novel human cancer-related genes and proteins (ABZ78120-ABZ78168 and 1789-ABR01681). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful as probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
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                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New composition comprising a substance that modulates the structure of proteins and polynucleotides, useful for therapeutic, prognostic and diagnostic reagents for eliciting cellular or humoral immune response:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200283921-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human leukocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human cancer-related protein 83P4B8 HLA peptide #1559.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AGEN-) AGENSYS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; vaccine;
                                                                                                                                                                  9
                                                                                                                                                                                                                    Similarity
7; Conserv
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                                                                                                                                                                ROOFIOSVLNNGATR 23
                                                                                                                                                                                                                                                                                                                                                                                             invention
                                                                                                           RQEILEQVLNRVVTR 15
                                                                                                                                                                                                                                                                                                                                  ₽,
                                                                                                                                                                                                                          Conservative
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2001US-0283112P.
2001US-0286630P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antigen
                                                                                                                                                                                                                                              29.8%;
46.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
                                                                                                                                                                                                                       Score 37; DB Pred. No. 75; 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₹
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ano AB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       response; HLA;
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                                                                                                                                                                                                                                                                           Length 15;
                                                                                                                                                                                                                          Indels
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RESULT 3
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Matches
                                                                                                                                                                                                                                                                                                The present invention relates to novel human cancer-related genes and proteins (ABZ78120-ABZ78168 and ABR01789-ABR01861). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful as probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the diagnosis and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcripts, as therapeutic agents. The proteins and peptides are useful as therapeutic, prognostic and diagnostic reagents for cancer. The present sequence is a human leukocyte antigen (HLA) peptide, used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New composition comprising a substance that modulates the structure of proteins and polynucleotides, useful for therapeutic, prognostic and diagnostic reagents for eliciting cellular or humoral immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-APR-2001; 2001US-0282739P
10-APR-2001; 2001US-0283112P
25-APR-2001; 2001US-0286630P
                                                                                                                                                                                                                                                                             Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                  HLA binding epitope #5247.
                                                           10-MAR-2005
                                                                                                             ADW34497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 13; Page 491; 1021pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer patients.
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Morrison K, 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human leukocyte antigen.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic; vaccine; cancer; immune response; HLA;
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                                                                                                                                                                                                                          Similarity 7; Conserv
                                                                                                                                                                                               ROOFIOSVLNNGATR
                                                                                                          standard; peptide; 13
                                                                                                                                                                           ROEILEOVLNRVVTR
                                                                                                                                                                                                                                                                              AA;
                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Morrison
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                                                           (first
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                                                           entry)
                                                                                                                                                                                                                                        29.8%;
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••
                                                                                                                                                                                                                                      Score 37;
Pred. No.
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                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                    DB .
75;
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Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T cell; MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA;

0,

viral disease; cancer.

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ADW35730
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AC ADW3
AC ADW3
AC ADW3
AC ADW3
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DT 10-M
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DE HLA
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KW Viru
KW WHC
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PD 15-M
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Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an ACL or estricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnosting viral diseases and cancer. The peptide epitopes are useful as diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
                                                                                              Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T cell; MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA; viral disease; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a composition comprising at least one peptide having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The
   15-MAY-2003
                                                                   Unidentified
                                                                                                                                                                HLA binding
                                                                                                                                                                                                  10-MAR-2005
                                                                                                                                                                                                                                                                ADW35730 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                binding motifs for HLA, us viral diseases and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing
                                WO2003040165-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 52-379; 382pp; English.
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20-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                   N
                                                                                                                                                                                                                                                                                                                                                                                                                   9; Conserv
                                                                                                                                                                                                                                                                                                                                                                    KYRQQFIQSVLNNGAT 22
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                                                                                                                                                                 epitope #6480
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                                                                                                                                                                                                (first entry)
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2001US-0285624P
                                                                                                                                                                                                                                                                                                                                                                                                                                27.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                   Score 34; DB
Pred. No. 2e+(
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                DB 7;
2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 13;
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RESULT 5
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Matches
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01-JUL-2002
08-OCT-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an AL restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allels-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnosing viral diseases and cancer. The peptide epitopes are useful as diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
                           11-JUL-1986;
10-JUL-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a composition comprising at least one peptide having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The
                                                                     10-JUL-1987;
                                                                                                                                                           Synthetic
                                                                                                                                                                                          oncogene related peptide; hst-related
                                                                                                     20-JAN-1988
                                                                                                                                                                                                                        hst-related
                                                                                                                                                                                                                                                                                                               AAP81118;
                                                                                                                                                                                                                                                                                                                                         AAP81118 standard; protein; 19 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 52-379; 382pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        viral diseases and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-441519/41.
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20-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   position comprising at least one peptide having allele-specific motifs for HLA, useful for preventing, treating or diagnosing
                                                                                                                                                                                                                                                                                                                                                                                                                                                 KYRQQFIQSVLNNGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sidney J,
                                                                                                                                                                                                                      peptide hst (169-187).
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                                                                                                                                                                                                                                                  (revised)
(revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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2001US-0285624P
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                           86JP-00164331.
87JP-00173671.
                                                                       87EP-00109985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27.4%;
56.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Southwood
                                                                                                                                                                                                                                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 34; DB 7;
Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; - Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 13;
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RESULT 6
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Best Local
Ine present invention relates to novel human cancer-related genes and proteins (ABZ78120-ABZ78168 and ABR01789-ABR01861). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful as probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the diagnosis and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcription.
                                                                                                                                                                                                                  Jakobovits A
Morrison K,
                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prepd by solid phase or soln. methods. One of 2 among 23 claimed oncogene-related peptides. See AAP81119-P81134. (Updated on 01-JUL-2002 to add
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                            Claim
                                                                                                                                            New composition comprising a substance that modulates the structure proteins and polynucleotides, useful for therapeutic, prognostic and diagnostic reagents for eliciting cellular or humoral immune respons
                                                                                                                                                                                                                                                                                      10-APR-2001;
10-APR-2001;
                                                                                                                                                                                                                                                                                                                                                 24-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                Human; cytostatic; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cancer-related protein 83P4B8 HLA peptide #1526.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR30464 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oncogene related peptide(s) - useful in prepn. of antigen and antibodies having specific reactivity with cancer proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Abe
                                                                                                                                                                                                                                                                                                                           10-APR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                              sapiens.
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25-APR-2001; :
                                                                                                                                                                     The present invention relates to novel human cancer-related genes and proteins (ABZ78120-ABZ78168 and ABR01789-ABR01861). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful as probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the diagnosis and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcripts, and inhibiting the expression of genes and/or translation of transcripts, and inhibiting the expression of genes and/or translation of transcripts, and inhibiting the expression of genes and/or translation of transcripts, and inhibiting the expression of genes and/or translation of transcripts, and inhibiting the expression of genes and/or translation of transcripts, and inhibiting the expression of genes and/or translation of transcripts.
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                                                                                  inhibiting the expression of genes and/or translation of transs as therapeutic agents. The proteins and peptides are useful as therapeutic, prognostic and diagnostic reagents for cancer. The sequence is a human leukocyte antigen (HLA) peptide, used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New composition comprising a substance that modulates proteins and polynucleotides, useful for therapeutic, diagnostic reagents for eliciting cellular or humoral
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leukocyte antigen.
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Pred. No.
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25-APR-2001;
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Morrison K, I
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                                                                                                                                                                                                                                                                                                                                                                       ; Score 32; DB; Pred. No. 4.9e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Faris M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ΑB;
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                                                                                                                                                                                                                                                                                                                                                           DB 6;
, 4.9e+02;
5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the structure of prognostic and immune response in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RS;
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RESULT 10 ABJ01432

ABJ01432 standard; peptide;

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19-SEP-2002

(first

entry)

Cytostatic; 158P1D7; cancer; bladder cancer;

mouse; rat; rabbit;

158P1D7 related HLA peptide

SEQ ID No 132.

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ESYKHEQVYIRS

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Best Loc
Matches
                                                                                                                    Query Match
                                                                                                                                                                                                                               diagnosing viral diseases and cancer. The peptide epitopes are useful as diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
                                                                                                                                                                                                                                                                                                                                                    composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an A2 restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a composition comprising at least one peptide having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytoroxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The
                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing viral diseases and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA; viral disease; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 52-379; 382pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-OCT-2000;
20-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HLA binding epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-OCT-2001; 2001WO-US051650.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-MAY-2003.
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                                                                                        Local
                                                        Similarity 5; Conserv
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QNYKYRQQFIQS 15
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                                                           Conservative
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2001US-0285624P
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                                                                                     25.8%;
41.7%;
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                                                        5.
                                                     Score 32; DB 7; L
Pred. No. 4.9e+02;
5; Mismatches 2;
                                                                                                                 Length 15;
                                                           Indels
                                                           0
                                                        Gaps
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Best Local
                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel nucleic acid, designated 158P1D7. The compositions are useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. mice, rats, rabbits, dogs, cate, cows, horses or humans). The compositions are also useful for monitoring genetic abnormalities and in preparing cancer vaccines. The nucleic acid of the invention can be used in gene therapy to treat the said disorders. This sequence represents a human leukocyte antigen (HLA) peptide relating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cat; cow; horse;
HLA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New compositions comprising a gene (designated 158P1D7), its encoded protein or their modulators, useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. dogs, cats, cows, horses
           22-AUG-2000;
10-APR-2001;
                                              22-AUG-2001; 2001WO-US026276
                                                                                               WO200216593-A2
                                                                                                                       Homo sapiens
                                                                                                                                                                       Cytostatic; 158P1D7; cancer; bladder cancer; mouse; rat; rabbit;
                                                                                                                                                                                               158P1D7
                                                                                                                                                                                                                      19-SEP-2002
                                                                                                                                                                                                                                                                   ABJ01921 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Faris M, Hul
Challita-Eid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AGEN-)
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10-APR-2001; 2001US-0282739P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-AUG-2001; 2001WO-US026276
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200216593-A2
                                                                                                                                                         cow; horse; human; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens.
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                                                                                                                                                                                                                                                                                                                                                                              Similarity 6; Conserv
                                                                                                                                                                                              related HLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                        SVLNNGAT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hubert RS, Raitano AB, Eid PM, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 130; 181pp; English
                                                                                                                                                                                                                                                                                                                                                                                Conservative
            2000US-0227098P
2001US-0282739P
                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human; vaccine; gene therapy; human leukocyte antigen;
                                                                                                                                                                                            peptide
                                                                                                                                                                                                                                                                                                                                                                                            25.0%;
75.0%;
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                                                                                                                                                                                              SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                          Score 31; DB 5;
Pred. No. 2e+06;
                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                         gene therapy; human leukocyte antigen;
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                                                                                                                                                                                                                                                                                                                                                                                                      Length 9;
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RESULT 12
ABJ01457
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel nucleic acid, designated 158P1D7. The compositions are useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. mice, rate, rabbits, dogs, cate, cows, horses or humans). The compositions are also useful for monitoring genetic abnormalities and in preparing cancer vaccines. The nucleic acid of the invention can be used in gene therapy to treat the said disorders. This sequence represents a human leukocyte antigen (HLA) peptide relating to the 158P1D7 protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New compositions comprising a gene (designated 158P1D7), its encoded protein or their modulators, useful for treating or diagnosing cancer particularly bladder cancer, in mammals (e.g. dogs, cats, cows, horse
                                                                                                                                                                                                                                                                                                                                                                                           158P1D7 related HLA peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 148; 181pp; English.
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Challita-Eid PM, Jakobovits
                                               New compositions comprising a gene (designated 158P1D7), its encoded protein or their modulators, useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. dogs, cats, cows, horses
                                                                                                                                                                                                                                                                                                                                                                 Cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                      19-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABJ01457
                                                                                                    WPI; 2002-425659/45.
                                                                                                                             Challita-Eid
                                                                                                                                          Faris M,
                                                                                                                                                                                             22-AUG-2000; 2000US-0227098P
10-APR-2001; 2001US-0282739P
                                                                                                                                                                                                                                    22-AUG-2001; 2001WO-US026276
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                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AGEN-) AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                      cow; horse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15
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                                                                                                                                                                     AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SVLNNGAT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SLLNNGLT 9
                                                                                                                                          Hubert RS,
                                                                                                                                                                                                                                                                                                                                                                   158P1D7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                              PM,
                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                              Jakobovits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25.0%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                     cancer; bladder cancer; mouse; rat; rabbit; do
an; vaccine; gene therapy; human leukocyte anti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Raitano AB,
bovits A;
                                                                                                                                          Raitano AB,
                                                                                                                                                                                                                                                                                                                                                                                            SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in mammals (e.g. dogs, cats, cows, horses
                                                                                                                              P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 31; DB 5;
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                            ID No
                                                                                                                                            Afar DEH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Afar
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                                                                                                                                            Levin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 10;
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Disclosure;

Page 131; 181pp; English

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RESULT 13
AAW19443
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
The present sequence represents peptide Q2.3.9.10, where at least one amino acid is in the D-configuration. The peptide preferably is modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The peptide, based on the naturally occurring histidinerich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic Candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and
                                                                                                                                                                                                                                                                                                                             Anti-fungal and anti-bacterial histatin-based peptide(s) treating Candidal infections, periodontitis and caries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     compositions are useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. mate, ratbits, dogs, cats, cows, horses or humans). The compositions are also useful for monitoring genetic abnormalities and in preparing cancer vaccines. The nucleic acid of the invention can be used in gene therapy to treat the said disorders. This sequence represents a human leukocyte antigen (HLA) peptide relating to the 158P1D7 protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-052232/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JUN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vaginal infection; urethral infection; mucosal infection; ea
respiratory infection; skin infection; ophthalmic infection;
bacterial disease; Streptococcus mutans; Porphyromonas gingi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Candida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW19443 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti-fungal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PERI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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he 158P1D7 protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ٠<u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            ଟି
                                                                                                                                                                                                                                                                              Fig 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and anti-bacterial histatin-based peptide 13-Q2.3.9.10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Roberts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 31; DB 5;
Pred. No. 4.4e+02;
1; Mismatches 1
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2 12

The present sequence represents an anti-fungal and anti-bacterial D-amin acid histatin-based peptide 113-Q2.3.9.10, found in human paratoid secretion, where at least one amino acid in the peptide is in the D-configuration. The peptide is preferably modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C- terminus. The movel D-amino acid-containing peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g.

Disclosure;

Fig 1; 63pp;

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                                                                                                                                                                                                                                                                                                                                                                                                                             vaginal infection; urethral infection; mucosal infection; respiratory infection; skin infection; ophthalmic infection bacterial disease; Streptococcus mutans; Porphyromonas gi
                                                                                                                                                           Anti-fungal and anti-bacterial D-amino acid histatins and histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis
                                                                                                                                                                                                             Oppenheim FG,
                                                                                                                                                                                                                                                                07-JUN-1995;
                                                                                                                                                                                                                                                                                     07-JUN-1996;
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                                                                                                                                                                                                                                                                                                                                                                   Region
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55.6%;
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Pred. No. 5.5e+02;
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Best Local
                            The invention relates to treating cystic fibrosis via the administration of a histatin, a fragment of histatin, or a histatin related peptide. The peptide that is administered contains between 8 and 20 amino acids and has the sequence: X1-X2-X3-X4-X5-X6-X7-X8-X9-X10-X11-X12-X13-X14-X15-X16-X17-X18-X19-X20-X21-X22-X23 where X1 is Asp or absent; X2 is Ser or absent; X3 is His or absent; X4 is Ala, His, Leu or absent; X5 is Lys, Gln, Arg, Orn, or another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X7 is His, Phe, Tyr Leu, or another hydrophobic amino acid; X8 is His, Phe, Tyr Leu, or another hydrophobic amino acid; X8 is His, Phe, Tyr Leu, or another hydrophobic amino acid; X10 is Tyr; X11 is Lys, His, Phe, Leu, or another hydrophobic amino acid; X10 is Lys, Gln, Arg, Orn, or another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X14 is Phe or absent X15 is His, Phe, Tyr,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Page 13; 27pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cystic fibrosis and combating Pseudomonas and other pulmonary ns in a mammal comprises administering histatin or a fragment
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   Phe or absent X15 is His, Phe, Tyr, or absent; X16 is Glu or absent; X17
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The invention relates to treating cystic fibrosis via the administration of a histatin, a fragment of histatin, or a histatin-related peptide. The peptide that is administered contains between 8 and 20 amino acids and has the sequence: X1-X2-X3-X4-X5-X6-X7-X8-X9-X10-X11-X12-X13-X14-X15-X16-X17-X18-X19-X20-X21-X22-X23 where X1 is App or absent; X2 is Ser or absent; X3 is His or absent; X4 is Ala, His, Leu or absent; X5 is Lys, Gln, Arg, Orn, or another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X7 is His, Phe, Tyr Leu, or another hydrophobic
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Synthetic.
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                                                                                                                                                                                                                                                                                     Example
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                                                                                                                                                                                                                                                                                     Page 12;
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Pred. No. 5
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5.5e+02;
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fragment
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RESULT 17
ABB56747
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Best Local
ABL00010 to ABL01104 represent human nucleic acid oligonucleotides comprising one or more single nucleotide polymorphisms (SNPs). ABB: to ABB56903 represent human peptides encoded by some of the SNP oligonucleotides. The sequences from the present invention can have
                                                                                                                                                                                                                              Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a pathology, e.g. autoimmune diseases, ascribed to the presence of a sequence
                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-355949/37.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-NOV-1999;
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immunosuppressive; antiinflammatory; neuroprotective; antimicrobial;
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                                                                                                                                               Page
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                                                                                                                                            636; 674pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid sequence SEQ ID NO:1312.
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55.6%;
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3; Mismatches
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5.5e+02;
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The present i recruitment d it. The CARD

invention relates to the isolation of human caspase domain-12 (CARD-12), and the polynucleotide sequence encoding D family of proteins are involved in apoptosis. The invention

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Page 18; 112pp; English.

Novel caspase recruitment domain-12, CARD-12, polypeptide useful for treating inflammatory disorders and immune disorders e.g., Crohn's disease, diabetes, multiple sclerosis, Hashimoto's thyroiditis and Grave's disease.

WPI; 2003-093098/08

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RESULT 18
ABG72213
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunosuppressive, cytostatic, antiinflammatory, neuroprotective and antimicrobial activities. Nucleic acids, polypeptides, oligonucleotides and antibodies from the present invention can be used for treating a subject suffering from, at risk for, or suspected of, suffering from a pathology ascribed to the presence of a sequence polymorphism. The pathology may be autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms. The SNPs and useful for determining which forms of a characterised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; caspase recruitment domain-12; CARD-12; apoptosis; transgenic; inflammatory disorder; immune disorder; Crohn's disease; reactive arthritis; insulin-dependent diabetes; multiple sclerosis;
                                                                                                                                                                                                                  Bertin
                                                                                                                                                                                                                                                                                  24-APR-2001; 2001US-00841739
                                                                                                                                                                                                                                                                                                                    24-APR-2002; 2002WO-US013009
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cerebroprotective; immunogen.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphism are present in individuals. The antibodies may be used in the detection, quantitation and/or cellular or tissue localisation of a polymorphic protein (e.g., for use in measuring levels of the polymorphic protein within appropriate physiological samples)
                                                                                                                                                                                                                                                   (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
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60.0%;
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Pred. No. 6.5e+02;
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Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  also discloses CARD-12 fusion proteins, antigenic peptides, anti-CARD-12 antibodies, recombinant expression vectors, host cells containing these expression vectors, and non-human transgenic animals. CARD-12 polypeptide and polynucleotide sequences are useful for diagnosing and treating inflammatory and immune disorders such as Crohn's disease, reactive arthritis, Lyme disease, insulin-dependent diabetes, multiple sclerosis, Hashimoto's thyroiditis, Grave's disease, contact dermatitis, psoriasis,
The invention relates to novel retro-inverso peptides or peptide-like molecules which have a high probability of binding to and/or otherwise modulating the function of polypeptides and/or proteins, and to method
                                                                                                                                                                      Novel isolated peptide comprising retro-inverso amino acid sequence of peptide, useful for diagnosing or treating cancer.
                                                                                                                                                                                                                                                           WPI; 2005-151359/16
                                                                                                                                                                                                                                                                                                                       Mandell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-FEB-2004; 2004US-00777829
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human NGF receptor targeted peptide SEQ ID NO:31, retro-inverso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-APR-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADW86333 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15
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peptide of human CARD-12 is
                                                                                                                                                                                                                                                                                                                                                                                                       (MAND/) MANDELL A J. (SELZ/) SELZ K A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nerve growth factor; cytostatic; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N
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                                                                                                                                                                                                                                                                                                                                                                            SHLESINGER M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         detection; screening; neoplasm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WRQESLOSVKN 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YROOFIOSVLN 18
                                                                                                                Page; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B
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2001US-00767460.
                                                                                                                                                                                                                                                                                                                 Selz KA,
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1. .15
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Pred. No. 7.1e+02;
3; Mismatches 2;
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  to methods
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RESULT 20
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ID AAW73692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            carried. A retro-inverso peptide of the invention is useful for diagnosing or treating cancer, or for detecting contaminants in water, food or soil samples. The present sequence represents a retro-inverso form of a peptide designed to target the human nerve growth factor (NGF) receptor. Note: The present sequence is not shown in the specification, but was derived from ADW86234.
                                                                                                                                                                                                               protective immunity in a patient against M. tuberculosis. Tuberculosis is a chronic, infectious disease generally caused by M. tuberculosis infection, and if left untreated typically results in serious complications and death. Fusion proteins containing the antigen, or DNA molecules can similarly be included with an acceptable carrier in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    M. tuberculosis antigen clone ORF-1 peptide 1-16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for designing such peptides or peptide-like molecules. Certain peptides of the invention have cytostatic activity, depending on their polypepti
                       response (especially induration) on the patient's skin. Inhibiting the spread of tuberculosis requires vaccination and accurate diagnosis, since antibiotic therapy may not be effective due to the existence of an
                                                                                  pharmaceutical compositions or in vaccines and administered as above. The polypeptides are also useful for diagnosis of tuberculosis, by contacting dermal cells with at least one polypeptide and detecting an immune response (especially induration) on the patient's skin. Inhibiting the
                                                                                                                                                                                                                                                                                                                                                                             tuberculosis antigen of the invention. The polypeptides are useful for immunotherapy to treat or prevent tuberculosis (especially in humans), e.g. they can be included with an acceptable carrier in pharmaceutical compositions or included in vaccines, and administered to induce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated Mycobacterium tuberculosis antigens - used products for the prevention, treatment and diagnosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAY-1997;
05-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence represents an immunogenic portion of a Mycobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 55; 104pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-045315/04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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98US-00073009
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Pred. No. 7.1e+02;
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RESULT 21
AAW73802
                                                                This sequence represents an immunogenic portion of a Mycobacterium cuberculosis antigen of the invention. The polypeptides are useful for immunotherapy to treat or prevent tuberculosis (especially in humans), ce e.g. they can be included with an acceptable carrier in pharmaceutical compositions or included in vaccines, and administered to induce protective immunity in a patient against M. tuberculosis. Tuberculosis is can fection, and if left untreated typically results in serious complications and death. Fusion proteins containing the antigen, or DNA complications and death. Fusion proteins containing the antigen, or DNA completed and included with an acceptable carrier in pharmaceutical compositions or in vaccines and administered as above. The playmate cells with at least one polypeptide and detecting an immune cesponse (especially induration) on the patient's skin. Inhibiting the spread of tuberculosis requires vaccination and accurate diagnosis, since casympetomatic but contagious stage and to patient on-compliance. The polypeptides overcome concerns of safety and efficacy of current vaccination with live bacteria (usually Bacillus Calmette-Guerin) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polypeptide comprising immunogenic Mycobacterium tuberculosis antigen useful for immunisation against M. tuberculosis infection to treat or prevent tuberculosis, and in diagnosis of tuberculosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-MAY-1997;
05-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptides overcome concerns of safety and efficacy of current vaccination with live bacteria (usually Bacillus Calmette-Guerin) lack of sensitivity and specificity of existing diagnostic technic
Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 54; 100pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  M. tuberculosis antigen clone ORF-1 peptide 1-16.
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                                                     lack of sensitivity and specificity of existing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CORI-) CORIXA
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Similarity 33.3%;
5; Conservation
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NYEQQEQASQQILSS 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dillon DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-00859381
98US-00073010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US010407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Skeiky YAW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 31; DB 2;
Pred. No. 7.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ⋛
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Campos-Neto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 16
                                                     diagnostic techniques
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RESULT 22
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Matches 5
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Best Local (
                                                                                                                                                                                    The present invention relates to the isolation of Mycobacterium tuberculosis antigen polypeptides (e.g. Tb224) and the nucleic acids encoding them. The invention describes compounds and methods for the diagnosis of tuberculosis or for inducing protective immunity against tuberculosis. The compounds comprise at least one immunogenic portion of one or more Mycobacterium proteins and nucleic acid molecules encoding such polypeptides. The Mycobacterium proteins and nucleic acid molecules encoding them can be used in diagnostic kits for the detection of Mycobacterium infection in patients and biological samples. The compounds of the invention and antibodies directed against the Mycobacterium proteins may be used in vaccines for immunisation against Mycobacterium proteins. The nucleic acids encoding the Mycobacterium proteins may be used in vaccines for immunisation against Mycobacterium proteins may be used in gene therapy. AMU08142-AMU08158 represent peptides (ORF-1-1 to ORF-1-17) which overlap to the open reading frame ORF-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 An isolated polypeptide comprising a Mycobacterium antigen, e.g., from Mycobacterium tuberculosis, useful in a vaccine for enhancing an immune
                                                                                                                                               Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Campos-Neto A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-FEB-2000; 2000US-0185037P 08-AUG-2000; 2000US-0223828P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 127; 161pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mycobacterium tuberculosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tuberculosis; Mycobacterium infection; gene therapy; anti bacterial; immunostimulant; mutant; mutein; ORF-1-16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mycobacterium tuberculosis overlapping peptide ORF-1-16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU08157;
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                                               Similarity
5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to and inhibiting development of a Mycobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NYEQQEQASQQILSS 16
NYKYROOFIOSVLNN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NYKYROOFIOSVLNN 19
                                                                                                                                               A
A
                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                        25.0%;
33.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25.0%;
  19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ovendale
                                               Score 31; DB
Pred. No. 7.6e
6; Mismatches
                                               6,
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Pred. No. 7.6e+02
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                                                                        .6e+02;
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                                                                                               Length 16
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                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                  Gaps
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RESULT 23

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ABG78004
ABG78004
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ABG
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                                                                                                                                                                                                                                                                                                                                                    with plasmid suspected of containing a DNA sequence that encodes a CD4+
CT cell stimulating antigen. Also included are (1) an isolated DNA
CS sequence consisting of DNA sequences isolated using the new method, their
CC complement or DNA sequences that hybridise to them, (2) an expression
CC vector comprising (1); (3) a host cell transformed with the expression
CC vector of (2); and (4) a polypeptide comprising an immunogenic portion of
CC antigen, or a variant of the antigen that differs only in conservative
CC substitutions and/or modifications, where the antigen comprises an amino
CC acid sequence encoded by (1). The method is useful for identifying and
CC isolating new antigens that stimulate CD4+ T cells. In particular, the
CC which the stimulation of CD4+ T cells is believed to play a role in the
CC which the stimulation of CD4+ T cells is believed to play a role in the
CC with infectious disease agents such as Mycobacterium tuberculosis or
CC is less technically difficult and less time-consuming than prior methods.
CC present method is also more suitable for high-throughput use. The
CC present sequence represents an overlapping peptide derived from a CD4+ T
                                                                                                                                  Matches
                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell stimulating antigens or antigens containing antibody epitopes comprises determining the level of CD4+ T cell stimulation when incubated with dendritic cells that have been exposed with a host cell transformed with dendritic cells that have been exposed with a best cell transformed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying DNAs encoding CD4+ T cell stimulating antigens, for isolanew CD4+ T cell-stimulating antigens associated with e.g. infections, comprises measuring the levels of interferon-gamma production or cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              M. tuberculosis CD4+ antigen ORF-1 1-1-1-17 peptide #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABG78004;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Skeiky YAW, Dillon DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     autoimmune
                                                                                                                                                                                                                                                                 Sequence 16
                                                                                                                                                                                                                                                                                                                               cell-stimulating antigen from M. tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to identifying DNA sequences that encode CD4+ T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-617730/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2002081579-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antigen; CD4+; infecautoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (POTT/)
N
                                                                 σ
                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Page 33; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POTTER J E R.
NYEQQEQASQQILSS 16
                                                                 NYKYROOFIOSVLNN 19
                                                                                                                                                                                                                                                                 AA;
                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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ease; Leishmaniasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-00023588
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                                                                                                                               Pred. No. 7.6e
6; Mismatches
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dendritic cell; T cel
                                                                                                                                                                 DB 5;
7.6e+02
                                                                                                                                                                                             Length 16;
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                                                                                                                               Gaps
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The invention provides a method for obtaining a plant which produces at cleast one seed having a modified phenotype. The method involves cransforming a host plant cell with a DNA construct which consists of transforming a host plant cell with a DNA construct which consists of coperably linked components in the direction of transcription, a promoter region from a Brassica plant gene, a DNA sequence of interest other than the native coding sequence, and a transcription termination region. The method is useful for obtaining plants having modified phenotype or for altering the phenotype of a plant seed or tissue. The DNA constructs are used in manipulating plant cells to provide for regulated transcription, cused in manipulating plant cells to provide for regulated transcription, constructs are also used as light inducible transcription, in a plant tissue or plant part of control. These constructs are also used for modulation of expression of control. These constructs are also used for modulation of expression of control. These constructs are also used for modulation products in the seed. Sequences AABS591-47 represent fragments of storage proteins used in comparison studies with the storage protein 2All
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-JUL-1990;
10-AUG-1993;
07-JUN-1995;
07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                         Obtaining a plant that produces a seed with a modified phenotype or altering a seed phenotype, comprises transforming a plant cell with construct consisting of operably linked components in the direction
Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-564354/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ricinus communis
                                                                                                                                                                                                                                                                                                        Example 9; Fig 6; 68pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Knauf VC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-MAR-1988;
29-APR-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JUL-1987;
25-JAN-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-JUL-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Castor bean
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-NOV-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CALJ ) CALGENE LLC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene; promoter; Brassica; phenotype; transcription; EA9; 3H11;
alpha-amylase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kridl JC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2S small subunit protein fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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90US-00526123.
90US-00550804.
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88US-00188361.
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                                                                                                                                                                                                                                                                                                                                                           of DNA
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QNLRQCQEYIQQV 14 ONYKYROOFIOSV 16 Matches Best Local Query Match

Similarity 6; Conserv

25.0%; 46.2%;

Score 31; DB 4; Pred. No. 8.2e+02; Mismatches

Length 17;

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Gaps

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Conservative

RESULT 24 AAB85946

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ADM37099
ID ADM3
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                                                                                                                                                                     CC and determining whether the candidate molecule causes a change in an CC expression of the label. Also included are a molecule capable of cc modulating activity of at least one kinase in situ identified by the CC claimed method, a fusion protein (comprising an E3 binding region, a CC kinase recognition domain and a green fluorescent protein, where the CC kinase recognition domain is the domain of beta-catenin, HIV protein VPU, p27, Bcl-2 or c-Jun), a fusion protein (comprising an E3 binding region, a kinase recognition domain and an enzyme capable of producing a CC detectable enzymatic product, where the kinase recognition domain is the domain of beta-catenin, HIV protein VPU, p27, Bcl-2 or c-Jun), an CC detectable enzymatic product, where the kinase recognition domain is the CC domain of beta-catenin, HIV protein VPU, p27, Bcl-2 or c-Jun), an CC vector capable of expressing the above genetic molecule and a cell cransfected with the above vector. Expression of the label requires CC phosphorylation of the phosphorylation substrate by the kinase. The CC whinse recognition domain is altered to include a consensus recognition CC whinse recognition on multiple kinase). The invention is useful to study kinase activities in situ and to screen for molecules that modulate kinase CC windows the kinase (e.g. AKT kinase). The invention is useful to study kinase activities in situ, for example in drug discovery. The invention allows for information on multiple kinases to be provided simultaneously, which conservation may be mutated to an AKT (not defined)
                                                                              Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying a molecule capable of modulating a kinase activity in situusing a substrate with an altered kinase recognition domain and an associated label is useful in drug discovery.
                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     kinase activity in situ, comprising exposing a candidate molecule to a cell comprising a phosphorylation substrate associated with a detectable label and having a kinase recogniston domain altered to be recognised by a kinase that does not recognise the substrate in its unaltered state,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
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US-08-331-224-60
US-08-331-223-60
US-08-331-231-59
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US-08-884-513-19
US-08-973-559-36
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ALIGNMENTS

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PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 157
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Patent No. 6863892
                                                                                                                                                      APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-BID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: OTHER CANCERS
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20050.00
CURRENT APPLICATION NUMBER: US/09/935,430
CURRENT FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
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PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
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APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20050.00
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CURRENT APPLICATION NUMBER: US/09/935,430
CURRENT FILING DATE: 2001-08-22
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APPLICANT: HUBERT, RE
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TYPE: PRT
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LEVIN, ELANA
CHALLITA-EID, PIA
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RAITANO, ARTHUR
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APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANTON: NUCLEIC ACID AND CORRESPONDING PRO
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETEC
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETEC
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-2005.00
CURRENT APPLICATION NUMBER: US/9/935,430
CURRENT APPLICATION NUMBER: 60/227,098
PRIOR PILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
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LENGTH: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RE
APPLICANT: RAITANO, A
                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
                                                                                        APPLICANT: Oppenheim, Frank G.
APPLICANT: Xu, Tao
APPLICANT: Spacciapoli, Peter
APPLICANT: Roberts, F. D.
APPLICANT: Friden, Philip M.
TITLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based
TITLE OF INVENTION: Deptides
NUMBER OF SEQUENCES: 37
                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook,
STREET: Two Militia Drive
STREET: Iwc
STREET: Lexington
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RAITANO, ARTHUR
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75.0%;
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75.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 31; DB
Pred. No. 79;
                                                         Smith & Reynolds,
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DETECTION OF BLADDER AND
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ZIP: 021/3 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy MEDIUM TYPE: IBM PC CON

Floppy disk

COUNTRY: US

Sala in the sale

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Patent No. 5912230
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Best Local Similarity 55.6%;
Matches 5; Conservative
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PACENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,563A
FILING DATE: 07-7UN-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/485,273
FILING DATE: 07-7UN-1995
ATTORNEY/AGENT INFORMATION:
NAME: NEOCK DAY/A F
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,559
FILING DATE: 07-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                           APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSE: Hamilton, Brook, Smith & Reynolds, P.0
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: PETELECOMMUNICATION INFORMATION:
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                                                                                                                                                               ZIP: 02173
                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                       STREET: Two Mil:
CITY: Lexington
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OTHER INFORMATION:
OTHER INFORMATION:
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TYPE: amino acid
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ROBERTS, F. D.
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must have a D configuration."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 6
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LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     Sequence 14, Application US/08505250 Patent No. 6183983
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Best Local Similarity
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Best Local Similarity 55.6%;
Matches 5; Conservative
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                                                                                                                                                                               APPLICANT: Sato, Haruya
APPLICANT: Yamamoto, Keiji
APPLICANT: Suzuki, Kokichi
APPLICANT: Ikeda, Masahiro
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APPLICANT: Rothstein, David M.
APPLICANT: Friden, Phillip M.
TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
CURRENT EPPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
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                                                                  APPLICANT: Sakagami, Masahiro
APPLICANT: Taniguchi, Makoto
TITLE OF INVENTION: PROTEIN MODIFICATION METHOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
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URRENT APPLICATION NUMBER: US/08/505,250
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LENGTH: 12 amino acids
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APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 QHHGYKQQF 11
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QHHGYKQQF 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25.0%;
55.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22,592
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Pred. No. 98;
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Pred. No. 98;
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; Sequence 49, Applicatic

; Patent No. 655563

; Patent No. FORMATION:

; GENERAL INFORMATION:

; APPLICANT: Aldersor

; APPLICANT: Skeiky.

; APPLICANT: Campos-
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EARLIER APPLICATION NUMBER: PCT/JP95/00298

EARLIER FILING DATE: 1995-02-27

EARLIER APPLICATION NUMBER: JP 198187/94

EARLIER FILING DATE: 1994-08-23

NUMBER OF SEQ ID NOS: 53

SOFTWARE: PATENTIN Ver. 2.0

SEQ ID NO 14

ENGTH: 14
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US-09-073-009-49
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                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: synthetic ; OTHER INFORMATION: peptide US-08-505-250-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-505-250-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 14
                                                                                                                                                                                                                                                                            Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: PCT/JP95/00298
PRIOR FILING DATE: 1995-02-27
PRIOR APPLICATION NUMBER: JP 198187/94
PRIOR FILING DATE: 1994-08-23
NUMBER OF SEQ ID NOS: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sakagami, Masahiro
APPLICANT: Taniguchi, Makoto
TITLE OF INVENTION: PROTEIN MODIFICATION METHOD
FILE REFERENCE: 110-511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/08/505,250 CURRENT FILING DATE: 1995-11-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sato, Haruya
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                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                          KKRSRFDQDVLN 14
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                                                                                                                                                                                                                        KYROOFIQSVLN 18
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                                                                                                    Application US/09073009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ikeda, Masahiro
Alderson, Mark
Dillon, Davin C.
Skeiky, Yasir A.W.
Campos-Neto, Antonio
                                                                                                                                                                                                                                                              Conservative
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58.3%;
                                                                                                                                                                                                                                                                              25.0%;
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Pred. No. 1.2e+02
                                                                                                                                                                                                                                                          Score 31; DB 2;
Pred. No. 1.2e+02
1; Mismatches
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                                                                                                                                                                                                                                                                                             Length 14;
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RESULT 10
US-09-073-010-49
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                                                                                                                                                                                                                                                                                      Patent No. 6613881

GENERAL INFORMATION:
APPLICANT: Alderson, Mark
APPLICANT: Dillon, Davin C.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Campos-Neto, Antonio
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS OF TUBERCULOSIS AND NUMBER OF SEQUENCES: 144
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 33.: Matches 5; Conservative
               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/073,010

FILING DATE: 05-MAY-1998

CLASSIFICATION:
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ORIGINAL SOURCE:
ORGANISM: Mycc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.441C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
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CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                   STREET: 6300 C
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                                                                                                                                                                                    ZIP: 98104
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                                                                                                                                                                                                   COUNTRY:
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33.3%;
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Pred. No. 1.4e+02;
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APPLICANT: Rodman, Toby C.
TITLE OF INVENTION: Monoclonal Human Natural Antibodies
FILE REFERENCE: 4436/12074-US1
CURRENT APPLICATION NUMBER: US/09/462,118
CURRENT FILING DATE: 1999-12-18
NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 30
LENGTH: 12
TYPE: PRT
ORGANISM: Human
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GENERAL INFORMATION:
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                        -09-623-548A-1034
                                                                               APPLICANT: Milner, Peter
APPLICANT: Holmes, Darren
APPLICANT: Thibaudeau, Karen
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD
TITLE OF INVENTION: COMPONENTS
                                                                                                                                                                                                APPLICANT: Conjuchem,
APPLICANT: Bridon, Do
CURRENT APPLICATION NUMBER: US/09/623,548A
CURRENT FILING DATE: 2000-09-05
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR FILING DATE: 1999-05-17
                                                                   FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 206-682-6031
INFORMATION FOR SEQ ID NO:
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ORIGINAL SOURCE:
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LENGTH: 16 amino acids
TYPE: amino acid
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5; Conservative
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                                                                                                                                                                                  Bridon, Dominique
Ezrin, Alan
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linear
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Pred. No. 1.4e+02;
6; Mismatches 4
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Pred. No. 1.4e+02;
3; Mismatches 1
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RESULT 14

PCT-US93-03748-1

Sequence 1, Application PC/TUS9303748 GENERAL INFORMATION:

APPLICANT: Howard, Russell J. APPLICANT: Leung, Lawrence L.K. TITLE OF INVENTION: Modulation

Modulation of Thrombospondin-CD36 Interactions

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                                                                                                                            ; ORGANISM: Artificial Sequence ; FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Synthetic ; OTHER INFORMATION: Peptide US-09-657-276-1034
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                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/13.
PRIOR FILING DATE: 1999-05-17
PRIOR APPLICATION NUMBER: 60/15:
PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: 60/15
PRIOR FILING DATE: 1999-10-18
                                                                                                                                                                                                                                                        SEQ ID NO 1034
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                                                                    Matches
                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Holmes, Darren
APPLICANT: Thibaudeau, Karen
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD
TITLE OF INVENTION: COMPONENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/657,276
CURRENT FILING DATE: 2000-09-07
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PRIOR APPLICATION NUMBER: 60/1
PRIOR FILING DATE: 1999-10-18
NUMBER OF SEQ ID NOS: 1617
                                                                                                                                                                                                                      LENGTH: 18
TYPE: PRT
                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE:
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                                                                 h 24.2%;
Similarity 41.7%;
5; Conservative
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AASHIYQNQFVQ 18
                                 AQNYKYRQQFIQ 14
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Pred. No. 2.3e
4; Mismatches
                                                               Score 30; DB 2; Pred. No. 2.3e+02;
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. 2.3e+02;
. 3;
                                                                                                  Length 18;
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US-08-816-105A-14
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Matches 5; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 5989882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 201-822-7039
TELEX: 219165
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                   TITLE OF INVENTION: ITTILE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: ITTILE OF INVENTION: UNMBER OF SEQUENCES:
                                                                                                         STREET: but
STREET: but
CITY: Tarrytown
CITY: New York
TISA
ZIP: 10591-5144
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4 Mb storage
COMPUTER: Apple Macintosh 6500
OPERATING SYSTEM: System 7.5
SOFTWARE: WordPerfect 3.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 30-APR-1992
ATTORNEY/AGENT INFORMATION:
ANAME: Dulak, NOTMAN C.
REGISTRATION NUMBER: 31,608
REFERENCE/DOCKET NUMBER: DX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRE
                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                               APPLICANT: Goeke, Kla
APPLICANT: Piepersber
APPLICANT: Stratmann,
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: O...
STREET: Madison
TTY: Madison
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh 6.0.5
SOFTWARE: Microsoft Word 4.00B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 07940
                                                                                                                                                                                 ADDRESSEE:
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                                                                                                                                                                E: Sprung Kramer Schaefer & Briscoe 660 White Plains Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 amino acids
                                                                                                                                                                                                                                                                                                                                          Crueger, Anneliese; Dellweg, Hans-Georg; Lenz,
Jurgen; Schroder, Werner; Pape, Hermann;
Goeke, Klaus; Schaper, Beate; Hemker, Michael;
Piepersberg, Wolfgang; Distler, Jurgen;
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                                                                                                                                                                                                                                      PROCESSES FOR PREPARING ACARVIOSYL TRANSFERASE AND FOR USING IT IN THE CONVERSION OF ACARBOSE HOMOLOGUES INTO ACARBOSE, FOR THE PREPARATION OF ACARBOSE HOMOLOGUES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24.2%;
41.7%;
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Pred. No. 2.3e+02;
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                                                                                                                                                    ; OTHER INFORMATION: Description of Artificial Sequence: Peptide US-09-935-430-180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: FARIS, MAI
APPLICANT: HUBERT, I
APPLICANT: RATTANO,
APPLICANT: AFAR, DAI
APPLICANT: LEVIN, E
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                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 180
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Best Local (
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APPLICATION NUMBER: DE 19625269.5
FILING DATE: 25-JUN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 19611252.4
FILING DATE: 22-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Kurt G. Briscoe
                                                                                                                                                                                                                                                                                                                                                             APPLICANT: AFAR, DANIEL
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: UUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20050.00
CURRENT FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
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INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                   LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (914) 332-1700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Kurt G. Briscoe
REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
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                                    12 FIQSVLNNG
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                                                                          Similarity 6; Conserv
  FQLSLLNNG
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Pred. No. 4.
                                                                                              Score 28; I
Pred. No. 2.
                                                                              Mismatches
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                                                                      2.5e+02;
2;
                                                                                                                  DB 2;
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PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PALENTIN Ver. 2.1
SEQ ID NO 425
LENGTH: 10
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US-09-411-706-6
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                                                                                                                                    ; OTHER INFORMATION: Description of Artificial Sequence: consensus ; OTHER INFORMATION: peptide US-09-411-706-6
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US-09-935-430-425
                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION: APPLICANT: Lane, K
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6, Application US/09411706 Patent No. 6312956
                                                                                                                                                                                                                                                        SEQ ID NO 6
                                                                    Matches
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                                                                                                 Query Match
                                                                                                                                                                                                                                                                      FILE REFERENCE: 3765
CURRENT APPLICATION NUMBER: US/09/411,706
CURRENT FILING DATE: 1999-10-01
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 51158-20050.00
CURRENT APPLICATION NUMBER: US/09/935,430
CURRENT FILING DATE: 2001-08-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: JAKOBOVITZ, AXA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION
TITLE OF INVENTION: OTHER CANCERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                  LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                     FEATURE:
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                                                                                  Similarity
                                 YKYRQQF 12
YKYRPEY 11
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RAITANO, ARTHUR
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                                                                 22.6%;
ilarity 57.1%;
Conservative
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                                                               Score 28; DB 2; LC
Pred. No. 3.1e+02;
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Pred. No. 2.
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                                                                                                 Length 12;
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                                                                   Indels
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OF BLADDER AND
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RESULT 19

US-07-918-181A-20

Application US/07918181A

Sequence 20, Patent No. 5

GENERAL INFORMATION

APPLICANT:

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US-09-543-940-9
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LENGTH: 12
TYPE: PRT
                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/128,106
PRIOR FILING DATE: 1999-04-07
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Huang, Ziwei
APPLICANT: Luo, Zhaowen
APPLICANT: Zhou, Naiming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 9, Apr-
Sequence 9, Apr-
No. 6613742
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APPLICANT: Lane, Kirk B.
TITLE OF INVENTION: NUCLEAR TARGETED PEPTIDE NUCLEIC ACID OLIGOMER
                                                                   Matches
                                                                                   Query Match
Best Local
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/935,032
CURRENT FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: US/09/411,706
PRIOR FILING DATE: 1999-10-01
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/543,940 CURRENT FILING DATE: 2000-04-06
                                                                                                                                                LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: peptide containing segment from C-terminal region
OTHER INFORMATION: of human SDF-1 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: consensus NLS OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                 PLICANT: Luo, Jiansong
                                                                                  Local
                              7 KYROQFIQSVLN 18
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                                                                                  Similarity
KWIQEYLEKALN 13
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                                                                   Conservative
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                                                                                  22.6%;
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Pred. No. 3.1e+02;
                                                                  Score 28; DB 2;
Pred. No. 3.4e+02
5; Mismatches
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                                                                                                Length 13;
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TITLE OF INVENTION:

C-Terminal IL-6 Muteins

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US-08-231-575-20
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Matches
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Patent No. 5565336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/231,575
FILING DATE: 22-APR-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/918,181
                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 212-645-2054
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                 APPLICANT: Fowlkes,
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: ImClone Systems Incorporated
STREET: 180 Varick Street
                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide HYPOTHETICAL: NO ANTI-SENSE: NO FRAGMENT TYPE: C-termin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Sheets, Eric J.
REGISTRATION NUMBER: 30,326
REFERENCE/DOCKET NUMBER: FO!
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 100
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/07/918,181A FILING DATE: 23-JUL-1992 CLASSIFICATION: 435
                                                                                                                                                                                          COUNTRY: U
                                                                                                                                                                                                                                      STREET: 180 Va. CITY: New York
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180 Varick Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 amino acids
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                                                                                                                                                                                                          United States
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23-JUL-1992
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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PCT-US93-06928-20
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Best Local 9
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INFORMATION FOR SEQ ID NO:
            MOLECULE TYPE: 1
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: (
                                                                                                                                                                                                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/918,181
FILING DATE: 23-JUL-1992
ATTORNEY/AGENT IMPORMATION:
ATTORNEY/AGENT IMPORMATION:
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HYPOTHETICAL: 1
ANTI-SENSE: NO
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                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 3
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                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: New York
STATE: New Yor
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nes 8; Conserv
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                                                                                                                                                                                                                              NAME: Feit, Irving N. REGISTRATION NUMBER:
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                                                                           TOPOLOGY:
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                                                                                      STRANDEDNESS:
                                                                                                                                                                                TELEPHONE:
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                                                                                                                                                                                                                             28,601
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Pred. No. 4e+02;
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; Seque...
; Patent No. 5670%;
; GENERAL INFORMATION:
; APPLICANT: Zhang, Shuguang
; APPLICANT: Lockehin, Curtis
; APPLICANT: Rich, Alexander
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Best Local Similarity
Thes 4; Conserva
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                                                                                                                                                                    Sequence 59, Appring No. 5670483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617) 861-9540
INFORMATION FOR SEQ ID NO: 58
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
TOPOLOGY: linear
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/973,326
FILING DATE: 28 DECEMBER 1992
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: MIT-6008
TELECOMMUNICATION INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 6: CORRESPONDENCE ADDRESS:
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APPLICANT: Lockshin, Curtis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
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CITY: Lexington
STATE: Massachu
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Pred. No. 4e+02;
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Search completed: January 26, Job time: 23.7184 secs

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Matches
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                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 861-6240
TELEFAX: (617) 861-9540
INFORMATION FOR SEQ ID NO: 59:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/346,84
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/973,326
FILING DATE: 28 DECEMBER 1992
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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TITLE OF INVENTION: STABLE MACROSCOPIC MEMBRANES FORMED BY
TITLE OF INVENTION: SELF-ASSEMBLY OF AMPHIPHILIC PEPTIDES AND USES
TITLE OF INVENTION: THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
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OPERATING SYSTEM:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A. ZIP: 02173-4799
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2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

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4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
     DMAQNYKYRQQFIQSVLNNGATRQ
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Copyright (c) 1993 - 2006 Compugen
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US-10-297-292-621
US-10-277-292-157
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ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/10/212,628
CURRENT FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 60/264,576
PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: US 60/305,232
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 10/057,789
PRIOR APPLICATION NUMBER: US 10/057,789
PRIOR FILING DATE: 2002-01-25
NUMBER OF SEQ ID NOS: 311
SOFTWARE: FASISEQ for Windows Version 4.0
SEQ ID NO 250
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US-10-057-789-250
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: DIFFERENTIAL LABELING FOR QUANTITATIVE TITLE OF INVENTION: ANALYSIS OF COMPLEX PROTEIN MIXTURES FILE REFERENCE: NADII.022A
CURRENT APPLICATION NUMBER: US/10/057,789
CURRENT FILING DATE: 2002-06-28
PRIOR APPLICATION NUMBER: US 60/264,576
PRIOR PILING DATE: 2001-01-26
PRIOR PILING DATE: 2001-07-13
NUMBER OF SEQ ID NOS: 311
CONTROLED OF SEG ID NOS: 311
                                                                                                                                                                                                                                                                                                                                            APPLICANT: Paul Haymes
APPLICANT: Jing Wei
APPLICANT: John Yates
APPLICANT: Nancy Andon
TITLE OF INVENTION: DIFFERENTIAL LABELING FOR QUANTITATIVE
TITLE OF INVENTION: ANALYSIS OF COMPLEX PROTEIN MIXTURES
FILE REFERENCE: NADII.022CP1
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APPLICANT: Jing Wei
APPLICANT: John Yates
APPLICANT: Nancy Andor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 17
TYPE: PRT
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FEATURE: NAME/KEY: VARIANT
                                   ORGANISM: Saccharomyces cerevisiae
                                                                   TYPE: PRT
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APPLICANT: AFAR, DANIEL
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20050.00
CURRENT APPLICATION NUMBER: US/9935,430
CURRENT FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTMARE: PATENTIN Ver. 2.1
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Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                  Sequence 132, Application US/10277292
Publication No. US20030199470A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 132
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Best Local
APPLICANT: AFAR, DANIEL
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: UCCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20050.00
CURRENT APPLICATION NUMBER: US/10/277,292
CURRENT FILLING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: US/09/935,430
PRIOR FILING DATE: 2001-08-22
                                                                                                                                                                                                                                                             APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: RAITANO, ARTHUR
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APPLICANT: HUBERT, RENE
APPLICANT: RAITANO, ARTHUR
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Pred. No. 3.6e+02;
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Pred. No. 1.7e+06
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                                                                                            US-10-990-137-132
                                                                                                                 RESULT 6
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US-10-280-340-132
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US-10-277-292-132
                                                     Sequence 132, Application US/10990137 Publication No. US20050227253A1
                                     GENERAL INFORMATION:
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Best Local
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NUMBER OF SEQ ID NOS: 700 '
SOFTWARE: Patentin Ver. 2.1
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CURRENT APPLICATION NUMBER: US/10/280,340
CURRENT FILING DATE: 2002-10-25
PRIOR APPLICATION NUMBER: US/09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
PRIOR FILING DATE: 2000-08-22
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APPLICANT: HUBERT, RENE
APPLICANT: RAITANO, ARTHUR
APPLICANT: FARIS, MARY APPLICANT: HUBERT, RE
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APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEPUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
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PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
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PRIOR FILING DATE: 2000-08-22
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ORGANISM: Artificial Sequence
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TYPE: PRT
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HUBERT, RENE
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75.0%;
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Pred. No. 1.7e+06;
1; Mismatches 1
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, OTHER INFORMATION: Description of Artificial Sequence: Peptide motif US-09-935-430-157
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US-09-935-430-157
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PRIOR ETILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 157
Matches
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                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                               APPLICANT: AFAR, DANIEL
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOUTZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20050.00
CURRENT APPLICATION NUMBER: US/09/935,430
CURRENT FILING DATE: 2001-08-22
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PRIOR FILING DATE: 2001-04-10
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
NUMBER OF SEQ ID NOS: 700
SOFTWARE: Patentin Ver. 2.1
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CURRENT APPLICATION NUMBER: US/10/990,137
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
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TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER ANT
TITLE OF INVENTION: OTHER CANCERS
                                                                                                                     LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
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b. US20030017466A1
Conservative
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                   25.0%;
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                 Score 31; DB 3;
Pred. No. 4.2e+02;

    Mismatches

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Pred. No. 1.7e+06;
Mismatches
1; Indels
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AFAR, DANIEL

APPLICANT: CHALLTA-EID, PIA

APPLICANT: JAKOBOVITZ, AYA

ITITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158PID7

ITITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND

TITLE OF INVENTION: OPHER CANCERS

FILTE OF INVENTION: OPHER CANCERS

FILTE REFERENCE: 51158-20050.00

CURRENT FILING DATE: 2001-08-22

PRIOR APPLICATION NUMBER: 60/227,098

PRIOR APPLICATION NUMBER: 60/227,098

PRIOR FILING DATE: 2000-08-22

PRIOR APPLICATION NUMBER: 60/227,098

PRIOR FILING DATE: 2001-08-20

PRIOR FILING DATE: 2001-08-20

PRIOR FILING DATE: 2001-08-21

PRIOR POLICATION NUMBER: 60/282,739

PRIOR POLICATION NUMBER: 60/282,739

PRIOR FILING DATE: 2001-04-10

NUMBER OF SEQ ID NOS: 700

SEQ ID NO 621

LENGTH: 10

TYPE: PRT

ORGANTS:
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GENERAL INFORMATION:
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 157
LENGTH: 10
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                                                           PRIOR APPLICATION NUMBER: US/09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
                                                                                                                                                                                                               APELICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: $1158-20050.00
CURRENT EPPLICATION NUMBER: US/10/277,292
CURRENT FILING DATE: 2002-10-21
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RAITANO, ARTHUR
AFAR, DANIEL
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CHALLITA-EID, PIA
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75.0%;
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Pred. No. 4.2e+02;
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APPLICANT: CHALLITA- EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20050.00
CURRENT APPLICATION NUMBER: US/01/277,292
CURRENT APPLICATION NUMBER: US/9935,430
PRIOR APPLICATION NUMBER: US/9935,430
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR FILING DATE: 2001-08-22
PRIOR FILING DATE: 2001-08-22
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 621
LENGTH: 10
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Best Local Similarity
Watches 6; Conserve
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US-10-280-340-157
; Sequence 157, Application US/10280340
; Publication No. US20030207835A1
; Publication No. US20030207835A1
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US-10-277-292-621
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GENERAL INFORMATION:

APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: AFAR, DANIEL
APPLICANT: AFAR, DANIEL
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOUTZ, AYA
APPLICANT: JAKOBOUTZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: OTHER CANCERS
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Publication No.
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Matches 6; Conserv
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ORGANISM: Artificial Sequence
FEATURE:
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RAITANO, ARTHUR
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Pred. No. 4.2e+02;
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Pred. No. 4.2e+02;
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CURRENT APPLICATION NUMBER: US/10/280,340 CURRENT FILING DATE: 2002-10-25

2002-10-25

FILE REFERENCE: 51158-20050.00

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RESULT 13
US-10-990-137-157
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; OTHER INFORMATION: Description of Artificial Sequence: Peptide motif
US-10-280-340-157
                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 621
LENGTH: 10
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US/09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
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NUMBER OF SEQ ID NOS: 700
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CURRENT APPLICATION NUMBER: US/10/280,340
CURRENT FILING DATE: 2002-10-25
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PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Peptide motif
                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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                                                                                                                                                                     Local Similarity
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LEVIN, ELANA
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75.0%;
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75.0%;
                                                                                                                                                                   Score 31; DB 4;
Pred. No. 4.2e+02;
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Pred. No. 4.2e+02;
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CURRENT APPLICATION NUMBER: US/10/990,137
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PACENTIN VET. 2.1
SEQ ID NO 621
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CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
PRIOR PILING DATE: 2001-04-10
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PACENTIN VEY: 2.11
SEQ ID NO 157
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APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEPTU IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: $11582005003
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APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION
TITLE OF INVENTION: OTHER CANCERS
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FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Peptide motif
                                              ORGANISM: Artificial Sequence
                                                                             LENGTH: 1
TYPE: PRT
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RAITANO, ARTHUR
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RAITANO, ARTHUR
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75.0%;
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Pred. No. 4.2e+02;
1; Mismatches 1
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OF BLADDER AND
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; NAME/KEY: VARIANT
; LOCATION: (7)...(0)
; OTHER INFORMATION: CSNP translation
US-10-813-638-1312
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US-09-226-666-14
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                                                                                                                                                                                                                                                                                                                                Sequence 1312, Application US/10813638

Publication No. US20040235026A1

GENERAL INFORMATION:

APPLICANT: Shimkets, Richard A.

APPLICANT: Leach, Martin D.

APPLICANT: Leach, Martin D.

TITLE OF INVENTION: NUCLEIC ACIDS CONTAINING SINGLE NUCLEIC ACID POLYMORPHISMS AND ME
TITLE OF INVENTION: USE THEREOF
FILE REFERENCE: 15966-599

CURRENT APPLICATION NUMBER: US/10/813,638

CURRENT FILING DATE: 2004-03-29

CURRENT FILING DATE: 2004-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14, Application US/09226666A Patent No. US20020077282A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/163,783
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1468
SOFTWARE: CuraGen Patent Formatter Version 0.9
SEQ ID NO 1312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 18
SOFTWARE: FRSTSEQ for Windows Version 4.0
SEQ ID NO 14
LENGTH: 12
TYPE: PRT
ORGANIZM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                     Query Match
Best Local :
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Best Local
                                    Matches
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CURRENT PEPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
                                                                                                                                                                                     LENGTH: 14
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                            FEATURE:
                                  Local Similarity 60.0
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les 6; Conserv
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 1 DMAQNYKYRQ 10
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55.6%;
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75.0%;
                                  Score 31; DB 5;
Pred. No. 6.1e+02;
0; Mismatches 4
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Pred. No.
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Pred. No. 5.1e+02;
3; Mismatches 1;
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                                                                   Length 14;
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US-09-073-009-49
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                                                                                                                                                                                            Sequence 49, Application US/09023588 Patent No. US20020081579A1
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Patent No. US2001001288BA1
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                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                               APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Dillon, Davin C.
APPLICANT: Alderson, Mark R.
TITLE OF INVENTION: METHOD FOR TH
NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 7
                                                                                                                                                           GENERAL INFORMATION: APPLICANT: Skeiky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 206-682-6031
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 211
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Dillon, Davin C.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Campos-Neto, Antonio
TITLE OF INVENTION: COMPOUNDS AN
NUMBER OF SEQUENCES: 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acida
STRANDENNESS: sin-
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,009
FILING DATE: 05-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
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     STREET: occurry: Seattle
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Washington
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Pred. No. 7e+02;
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                                      701 Fifth Avenue
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Query Match
Best Local Similarity
Thes 5; Conserve
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                                                                       JS-09-793-306-49
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Sequence 49, Mpp
No. US200
                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 49
 Matches
                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/185,037
PRIOR TILING DATE: 2000-02-25
PRIOR APPLICATION NUMBER: US 60/223,828
PRIOR FILING DATE: 2000-08-08
NUMBER OF SEQ ID NOS: 164
                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Compounds and Methods for Diagnosis and Immunotherapy TITLE OF INVENTION: of Tuberculosis FILE REFERENCE: 014058-008740US
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Campos-Neto, Antonio
APPLICANT: Skeiky, Yasir
APPLICANT: Ovendale, Pamela
                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/793,306
CURRENT FILING DATE: 2001-02-26
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: .
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acide
TYPE: amino acid
                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                       OTHER INFORMATION: Description of Artificial Sequence:peptide OTHER INFORMATION: ORF-1-16
                                                                                                                        FEATURE:
                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
ORIGINAL SOURCE:
ORGANISM: MYC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
Local Similarity nes 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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Corixa Corporation
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   Conservative
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                  25.0%; Score 31; DB 3; 33.3%; Pred. No. 7e+02;
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Pred. No. 7e+02;
   Mismatches
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                                    Length 16;
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US-10-251-526-30
Sequence 30, Application US/10251526
Publication No. US20030144486A1
GENERAL INFORMATION:
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US-10-936-237-11
                                                                                                                                                                    RESULT 22
                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Human US-10-247-946-30
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US-10-247-946-30
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Matches
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SEQ ID NO 11
LENGTH: 10
TYPE: PRT
ORGANISM: SARS Coronavirus
                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 30
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 30, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                         Query Match
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Matches 5; Conserv
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Publication No. US20050106563A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
APPLICANT: Rodman, Toby C.
TITLE OF INVENTION: Monoclonal Human Natural Antibodies
FILE REFERENCE: 4436/1C074-US1
CURRENT APPLICATION NUMBER: US/10/251,526
CURRENT FILING DATE: 2002-09-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 4436/1C074-US1
CURRENT APPLICATION NUMBER: US/10/247,946
CURRENT FILING DATE: 2002-09-20
PRIOR APPLICATION NUMBER: US/09/462,118
PRIOR FILING DATE: 1999-12-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Rodman, Toby C.
TITLE OF INVENTION: Monoclonal Human Natural Antibodies
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CURRENT FILING DATE: 2004-09-08
NUMBER OF SEQ ID NOS: 195
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APPLICANT: CHEN, LEE-HSUAN
TITLE OF INVENTION: EPITOPE PROFILE OF SARS CORONAVIRUS
FILE REFERENCE: 09468.0004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
                                                                                                                                                                                                                                                                                                Local Similarity tes 5; Conserv
                                                                                                                                                                                                                                                               2 MAQNYKYRQ 10
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No. US20030144485A1
                                                                                                                                                                                                                                                                                                    Conservative
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Pred. No. 7.4e+02;
3; Mismatches 1
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Pred. No. 6e+02;
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APPLICANT: BAKER, Matthew
APPLICANT: CARTER, Francis J.
APPLICANT: CARTER, Graham
ITILE OF INVENTION: METHOD FOR MAPPING AND ELIMINATING
ITITLE OF INVENTION: T-CELL EPITOPES
FILE REFERENCE: MER-135
CURRENT APPLICATION NUMBER: US/11/009,460
CURRENT FILING DATE: 2004-12-10
PRIOR APPLICATION NUMBER: PCT/EP03/06110
PRIOR APPLICATION NUMBER: EP02012919.3
                      GENERAL INFORMATION:

APPLICANT: K'ster, Hubert
APPLICANT: Siddiq1, Suhaib
APPLICANT: Siddiq1, Suhaib
APPLICANT: Siddiq1, Suhaib
APPLICANT: Little, Daniel
TITLE OF INVENTION: Capture Compounds, Collections
TITLE OF INVENTION: Compositions
TITLE OF INVENTION: Compositions
FILE REFERENCE: 24743-2305
CURRENT APPLICATION NUMBER: US/10/197,954
CURRENT FILING DATE: 2002-07-16
PRIOR APPLICATION NUMBER: 60/306,019
PRIOR APPLICATION NUMBER: 60/314,123
PRIOR FILING DATE: 2001-08-21
PRIOR FILING DATE: 2001-08-21
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US-11-009-460-104
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US-10-197-954-35
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US-10-251-526-30
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NUMBER OF SEQ ID NOS: 36
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 30
LENGTH: 12
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Matches 5; Conserv
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Publication No. US20030119021A1
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res 7; Conservarium
APPLICATION NUMBER: 60/363,433
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Pred. No. 7.4e+02;
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Query Match
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Marches 5; Conserve
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; NUMBER OF SEQ ID NOS: 149
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 35
; LENGTH: 18
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-197-954-35
Search completed: January Job time: 79 secs
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                                                                                                                                                                                                                                          TITLE OF INVENTION: Capture Compounds, Collections Thereof
TITLE OF INVENTION: And Methods For Analyzing The Proteome And Complex
TITLE OF INVENTION: Compositions
FILE REFERENCE: 24743-2309
CURRENT APPLICATION NUMBER: US/10/760,085
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: 60/441,398
PRIOR APPLICATION NUMBER: 60/441,398
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 149
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 35
LENGTH: 18
TYPE: PRT
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Best Local (
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Publication No. US20050042771A1
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APPLICANT:
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                                                                                                    3 AQNYKYRQQFIQ 14
                                                                   AASHIYQNQFVQ 18
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Subramaniam Marappan
Chester Frederick Hassman
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Pred. No. 1.2e+03;
4; Mismatches 3
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Pred. No. 1.2e+03;
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1: /cgn2_6/ptodata/1/pubpaa//USO8_NEW_PUB.pep:*

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3: /cgn2_6/ptodata/1/pubpaa//USO7_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa//PCT_NEW_PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa//USO7_NEW_PUB.pep:*

6: /cgn2_6/ptodata/1/pubpaa//USO7_NEW_PUB.pep:*

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US-11-111-463-12
US-11-045-024-13293
US-11-012-562-44
US-11-022-562-44
US-11-045-024-13006
US-11-045-024-13016
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Sequence 1123, Apsequence 255, Appl Sequence 75, Appl Sequence 9528, Apsequence 9528, Apsequence 9528, Apsequence 12, Appl Sequence 12, Appl Sequence 12, Appl Sequence 14, Appl Sequence 114, Appl Sequence 114, Appl Sequence 13006, Assequence 13016, Assequence 13016, Assequence 1215, Appl Sequence 215, Appl Sequence 215, Appl Sequence 229, Appl Sequence 229, Appl Sequence 239, Appl Sequence 239, Appl Sequence 256, Appl Sequence 283, Appl Sequence 256, Appl Sequence 256, Appl
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US-11-1045-024-1387
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US-11-045-024-13369
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US-11-053-039-123
US-11-054-515-229
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US-11-033-039-1123
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                                                                       CURRENT APPLICATION NUMBER: US/10/511,559
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR FILING DATE: 2002-04-18
PRIOR APPLICATION NUMBER: EP 03006554.4
PRIOR APPLICATION NUMBER: EP 03006554.4
PRIOR FILING DATE: 2003-03-24
NUMBER OF SEQ ID NOS: 1147
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-11-033-039-1123
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CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 10/245,871
PRIOR FILING DATE: 2002-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 10/197,000 PRIOR FILING DATE: 2002-07-17 PRIOR APPLICATION NUMBER: 09/396,813 PRIOR FILING DATE: 1999-09-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN
TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01
                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: MODIFIED FACTOR VIII FILE REFERENCE: MER-133
                                   LENGTH: 13
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                               APPLICANT: JONES
APPLICANT: BAKE
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LOCATION: (5)..(5)
OTHER INFORMATION:
ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: OTHER INFORMATION:
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Similarity 75.0%;
6; Conservation
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US-11-166-412-154
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Pred. No. 17;
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Sequence 154, App
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US-10-511-559-75
; Sequence 75, Application US/10511559
; Publication No. US20050256304A1
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-503-575-110
                                                                                                                                                                                                                                                                                                                                                                                                    US-10-503-575-110
                                                          NUMBER OF SEQ ID NOS: 340
SOFTWARE: PatentIn version 3.1
SEQ ID NO 110
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                                                                                                                     APPLICANT: KONING, FYILE

TITLE OF INVENTION: NOVEL EPITOPES FOR CELIAC DISEASE AND AUTOIMMUNE DISEASES, METHOD

TITLE OF INVENTION: DETECTING THOSE AND NOVEL NON-ANTIGENIC FOOD COMPOUNDS

FILE REFERENCE: 2799/72843-PCT-US

CURRENT APPLICATION NUMBER: US/10/503,575

CURRENT FILING DATE: 2004-08-04

PRIOR APPLICATION NUMBER: PCT/NL03/00077

PRIOR PILING DATE: 2002-02-04

PRIOR PILING DATE: 2002-02-04

PRIOR FILING DATE: 2002-02-04

PRIOR FILING DATE: 2002-02-04
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CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR APPLICATION NUMBER: EP 03006554.4
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APPLICANT: van Veelen, Petrus A
APPLICANT: Koning, Frits
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Pred. No.
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; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
US-11-045-024-2254
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CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR PRILING DATE: 1993-03-05
PRIOR PELLING DATE: 1993-03-05
PRIOR PELLING DATE: 1993-06-04
PRIOR PELLING DATE: 1993-06-04
PRIOR PELLING DATE: 1993-08-06
PRIOR PELLING DATE: 1993-08-06
PRIOR PELLING DATE: 1993-08-06
PRIOR PELLING DATE: 1993-10-29
PRIOR PELLING DATE: 1993-11-29
                                                                                                           Sequence 9528, Application US/11045024 Publication No. US20050271676A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PastSEQ for Windows Version 4.0 SEQ ID NO 2254
                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Epimmune Inc.
TITLE OF INVENTION: Inducing Cellular Responses to Human
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Ac
FILE REFERENCE: 2060.0040007
                                                       APPLICANT:
                                                                         APPLICANT: Sette, Alessandro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1994-12-01
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APPLICATION NUMBER: US 08/347,610
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Livingston, Brian
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Livingston, Brian
Chesnut, Robert
                                                   Sidney, John
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Grey, Howard M.
                                   Southwood, Scott
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sidney, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Robert
                                                                                                                                                                                                                                                                                                                     22.6%;
45.5%;
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Pred. No. 62
                                                                                                                                                                                                                                                                                                                       Score 28; DB 7;
Pred. No. 58;
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APPLICANT: Merzou APPLICANT: Arab, APPLICANT: Saxena APPLICANT: Eaves,

Saxena, Geeta Eaves, Connie J. Clark-Lewis, Ian

Merzouk, Ahmed

Lakhdar Christopher

APPLICANT:

APPLICANT: Clark-Lewis, Ian
APPLICANT: Salari, Hassan
APPLICANT: Chemokine Therapeutics Corporation
APPLICANT: The University of British Columbia
TITLE OF INVENTION: CXCR4 Antagonist Treatment
FILE REFERENCE: 080420-00010005

o f

Hematopoietic Cells

CURRENT APPLICATION NUMBER: US/10/945,674A
CURRENT FILING DATE: 2004-09-20
PRIOR APPLICATION NUMBER: CA 2,305,787
PRIOR FILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/205,467
PRIOR FILING DATE: 2000-05-19
PRIOR FILING DATE: 2000-05-19
PRIOR APPLICATION NUMBER: US 09/852,424
PRIOR FILING DATE: 2001-05-09
NUMBER OF SEQ ID NOS: 140
SOFTWARE: Patentin Ver: 2.0

SEQ ID NO 77 LENGTH: 13 TYPE: PRT

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US-10-945-674A-77
                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS US-11-045-024-9528
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TITLE OF INVENTION: Urus-1 Using Peptide a FILE REFERENCE: 2060.0040007
TITLE OF INVENTION: Virus-1 Using Peptide a FILE REFERENCE: 2060.0040007
CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-08-06
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,399
PRIOR FILING DATE: 1993-11-29
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILING DATE: 1993-11-29
PRIOR FILING DATE: 1993-11-29
GENERAL INFORMATION:
                    Sequence 77, Application US/10945674A
Publication No. US20060014682A1
                                                                                                                                                                                                                                                                                            Best Local Similarity 45.5%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 9528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
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Celis, Esteban
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                                                                                                                                                                                                                                                                                                             4; Mismatches
                                                                                                                                                                                                                                                                                                                                       Score 28; DB
Pred. No. 58;
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; NAME/KEY: MOD_RES; LOCATION: (1)
; OTHER INFORMATION: X
; OTHER INFORMATION: C;
; OTHER INFORMATION: 7
US-10-945-674A-77
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                                                                                                                                                                         ; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS US-11-045-024-13293
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US-11-045-024-13293
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CURRENT FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-06-04
PRIOR FILING DATE: 1993-08-06
PRIOR PRIOR DATE: 1993-08-06
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR FILING DATE: 1993-01-29
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1993-11-29
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR PRIOR DATE: 1994-03-04
                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL
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Publication No. US20050271676A1
                                                                                       Matches
                                                                                                         Query Match
Best Local
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Best Local :
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions FILE REFERENCE: 2060.0040007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Livingston, Brian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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OTHER INFORMATION:
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                                                                                                                                                                                                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION
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                                                                                       Similarity 5; Conserv
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4; Conservative
                                         QFIQSVLNNGA 21
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  QLIEALLDTGA 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Epimmune Inc.
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Grey, Howard M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Livingston, Brian
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                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Esteban
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Description of Artificial Sequence:synthetic SDF-1 CXCR4 receptor antagonist analogue C-terminal residues 55-67 linked by -(CH-2)-n-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Xaa = Leu linked via -(CH-2)-n-, where n =
to the C-terminal amino acid of any one of
78, 87-94, 103-110, and 117-122
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45.5%;
                                                                                       4; Mismatches
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Pred. No. 70;
                                                                                                       Score 28; DB
Pred. No. 82;
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SEQ ID NOS:76,
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TITLE OF INVENTION: MODIFIED FACTOR VIII
FILE REFERENCE: MER-133
CURRENT APPLICATION NUMBER: US/10/511,559
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT_P03/04063
PRIOR FILING DATE: 2003-04-17
PRIOR FILING DATE: 2002-04-18
PRIOR FILING DATE: 2002-04-18
PRIOR FILING DATE: 2003-04-17
PRIOR FILING DATE: 2003-04-18
PRIOR FILING DATE: 2003-03-24
NUMBER OF SEQ ID NOS: 1147
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 254
LENGTH: 13
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                                                                                                                                                  US-10-511-559-254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 254, Application US/10511559 Publication No. US20050256304A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 12
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                                                                                           Best
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Best Local
                                                                         Matches
                                                                                                           Query Match
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APPLICANT: BAKER, Matthew
APPLICANT: CARR, Francis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/11/111,463
CURRENT FILING DATE: 2005-04-21
PRIOR APPLICATION NUMBER: 60/564,313
PRIOR FILING DATE: 2004-04-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Turner, Jerrold R.
APPLICANT: Mrsny, Randall J.
APPLICANT: McKay, Derek
TITLE OF INVENTION: MYOSIN LIGHT CHAIN KINASE INHIBITORS AND METHODS OF USE
FILE REFERENCE: 092234-9030-US01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                    OTHER INFORMATION: Potential Epitope
                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                              TYPE: PRT
                                                                                                                                                                                        FEATURE:
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                                                                       Local Similarity hes 6; Conserv
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                                   YROOFIOSVLNNG 20
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                                                                         Conservative
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                                                                                           46.2%;
                                                                                                             21.8%;
                                                                       Score 27; DB 6;
Pred. No. 1e+02;
2; Mismatches
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Pred. No. 5.8e+04;
3; Mismatches 1
                                                                                                                                                                    of human
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                                                                                                           Length 13;
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RESULT 11

US-11-022-562-43

Sequence 43, Application US/11022562 Publication No. US20050249742A1

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RESULT 13
US-11-106-932-114
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CURRENT APPLICATION NUMBER: US/11/022,562
CURRENT FILING DATE: 2004-12-22
PRIOR APPLICATION NUMBER: PCT/US03/20322
PRIOR FILING DATE: 2003-06-27
PRIOR APPLICATION NUMBER: 60/392718
PRIOR FILING DATE: 2002-06-27
PRIOR FILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 340
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                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Human Immunodeficiency Virus US-11-022-562-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-11-022-562-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-11-022-562-43
                                                        Sequence 114, Application US/11106932 Publication No. US20050260697A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 44, Application US/11022562 Publication No. US20050249742A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FASTSEQ for Windows Version 4.0 SEQ ID NO 43
                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                           Matches
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING
TITLE OF INVENTION: A CYTOTOXIC T LYMPHOCYTE IMMUNE RESPONS
FILE REPERENCE: DFN-043CN
CURRENT APPLICATION NUMBER: US/11/022,562
CURRENT FILING DATE: 2004-12-22
FRIOR APPLICATION NUMBER: PCT/US03/20322
PRIOR APPLICATION NUMBER: PCT/US03/20322
PRIOR FILING DATE: 2003-06-27
APPLICANT: WANG, KA-WANG KEVIN
APPLICANT: HAYES, RONALD
APPLICANT: LIU, MING CHEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ruprecht, Ruth M.
APPLICANT: Shisong, Jiang
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING
TITLE OF INVENTION: A CYTOTOXIC T LYMPHOCYTE IMMUNE RESPONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/392718 PRIOR FILING DATE: 2002-06-27 NUMBER OF SEQ ID NOS: 340
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TYPE: PRT
ORGANISM: Human Immunodeficiency Virus
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                                                                                                                                                                                                                                                                                                               Local
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Pred. No. 1.2e+02;
1; Mismatches 3
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Pred. No. 1.2e+02;
1; Mismatches 3
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                                                                                                            ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS US-11-045-024-13006
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US-11-106-932-114
                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13006
                                   Best Local
Matches
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CURRENT FILING DATE: 2005-04-15
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PatentIn version 3.2
SEQ ID NO 114
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Best Local :
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PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2060.0040007
CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
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TITLE OF INVENTION: PROTEOLYTIC MARKERS AS DIAGNOSTIC BIOMARKERS FOR CANCER,
TITLE OF INVENTION: INJURY AND MUSCLE REHABILITATION/EXERCISE OVERTRAINING
FILE REFERENCE: 5853-549-1
                                                                                                                                                    TYPE: PRT
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                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 08/347,610
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TYPE: PRT
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APPLICATION NUMBER: US 08/205,713
FILING DATE: 1994-03-04
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15 SVLNNGATRO 24
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Similarity 60.0%;
6; Conservation
                                     Similarity 6; Conserv
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Chesnut, Robert
Baker, Denise Marie
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                                     Conservative
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                                                   21.8%;
                                   Score 27; DB 7; L
Pred. No. 1.2e+02;
1; Mismatches 3;
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Pred. No. 1.2e+02;
1; Mismatches 3
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US-11-045-024-13016

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                                                                                                                                        Sequence 8, Application US/10973977

Publication No. US20060008467A1

GENERAL INFORMATION:
APPLICANT: HAYNES, BARTON F.
APPLICANT: LIAO, HUA-XIN
APPLICANT: LETVIN, NORMAN
TITLE OF INVENTION: HUMAN IMMUNODEFICIENCY VIRUS VACCINE
FILE REFERENCE: 1579-942
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LENGTH: 15
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             CURRENT APPLICATION NUMBER: US/10/973,977
CURRENT FILING DATE: 2004-10-27
PRIOR APPLICATION NUMBER: 09/775,805
PRIOR FILING DATE: 2001-02-05
PRIOR APPLICATION NUMBER: 09/497,497
PRIOR FILING DATE: 2000-02-04
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NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
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NUMBER OF SEQ ID NOS: 107
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TYPE: PRT
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Local Similarity 60.0%;
les 6; Conservarium
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APPLICATION NUMBER: US 08/103,396
FILING DATE: 1993-08-06
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Celis, Esteban
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1; Mismatches
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US-11-145-861-215
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US-11-145-861-215
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LENGTH: 17
                                                                                                                                                                  SOFTWARE: PatentIn version 3.3 SEQ ID NO 1125
                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1125, Application US/11033039 Publication No. US20060002947A1
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Matches 5; Conserv
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TITLE OF INVENTION: Phage Microarray Profiling of the Humoral Response to Disease
FILE REFERENCE: UM-09899
CURRENT APPLICATION NUMBER: US/11/145,861
CURRENT FILING DATE: 2005-06-06
                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 10/245.871
PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-07-17
PRIOR PRIOR PRIOR DATE: 2002-07-17
PRIOR APPLICATION NUMBER: 09/396,813
PRIOR FILING DATE: 1999-09-14
                                                                                                                                                                                                                                                                                                                                   APPLICANT: XU, MINZHEN
TITLE OF INVENTION: L1-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01
CURRENT APPLICATION NUMBER: US/11/033,039
CURRENT FILING DATE: 2005-01-11
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SOFTWARE: PatentIn version 3.3
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                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 1452
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                                                    OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: hybrid peptide
                                                                                         ORGANISM: Artificial Sequence FEATURE:
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NAME/KEY: MOD_RES
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Pred. No. 1.4e+02
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Pred. No. 1.3e+02;
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Best Local Similarity
Warrches 6; Conserva
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US-11-022-562-249
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US-11-033-039-1125
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                                                                                  US-11-022-562-250
                                                                                                                        PRIOR APPLICATION NUMBER: 60/392718
PRIOR FILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 340
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 250
LENGTH: 20
TYPE: PRT
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 249
LENGTH: 20
                     Query Match
Best Local
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                                                                                                                                                                                                                                                            APPLICANT: Ruprecht, Ruth M.
APPLICANT: Shisong, Jiang
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING
TITLE OF INVENTION: A CYTOTOXIC T LYMPHOCYTE IMMUNE RESPONSE
FILE REFERENCE: DFN-043CN
FULL REFERENCE: DFN-043CN
CURRENT APPLICATION NUMBER: US/11/022,562
CURRENT APPLICATION NUMBER: US/11/022,562
CURRENT FILING DATE: 2004-12-22
PRIOR APPLICATION NUMBER: PCT/US03/20322
PRIOR FILING DATE: 2003-06-27
PRIOR FILING DATE: 2003-06-27
PRIOR FILING DATE: 2003-06-27
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING
TITLE OF INVENTION: A COMPOSITION T LYMPHOCYTE IMMUNE RESPONSE
FILE REFERENCE: DFN-043CN
CURRENT APPLICATION NUMBER: US/11/022,562
CURRENT APPLICATION NUMBER: US/11/022,562
CURRENT FILING DATE: 2004-12-22
PRIOR APPLICATION NUMBER: PCT/US03/20322
PRIOR PILING DATE: 2003-06-27
PRIOR APPLICATION NUMBER: 6/392718
PRIOR APPLICATION NUMBER: 60/392718
PRIOR FILING DATE: 2002-06-27
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                                                                                                     ORGANISM: Human Immunodeficiency Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
  Local Similarity 60.
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5; Conserv
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21.8%; Score 27; DB 7;
60.0%; Pred. No. 1.6e+02;
vative 1; Mismatches 3
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38.5%;
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Pred. No. 1.5e+02;
Pred. No. 5;
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Pred. No. 1.6e+02;
1; Mismatches
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3; Indels
                                        Length 20;
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; FEATURE:
; OTHER INFORMATION: Synthetic Peptide
US-11-196-400-16:
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; LENGTH: 19
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-503-575-329
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             Query Match
Best Local Similarity
                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 29
SOFTWARE: PatentIn version 3.3
SEQ ID NO 16
LENGTH: 20
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APPLICANT: DRUILHE, PIERRE
APPLICANT: DAUBERSIES, PI
Matches
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Matches 6; Conserv
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APPLICANT: KONING, Frits
TITLE OF INVENTION: NOVEL EPITOPES FOR CELIAC DISEASE AND AUTOIMMUNE DISEASES,
TITLE OF INVENTION: DETECTING THOSE AND NOVEL NON-ANTIGENIC FOOD COMPOUNDS
FILE REFERENCE: 2799/72843-PCT-US
CURRENT FILLING DATE: 2004-08-04
PRIOR APPLICATION NUMBER: PCT/NL03/00077
PRIOR APPLICATION NUMBER: PCT/NL03/00077
PRIOR FILING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: EP 02075456.0
PRIOR FILING DATE: 2002-02-04
NUMBER OF SEO ID NOS: 340
SOFTWARE: Patentin version 3.1
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                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US/09/742,096
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: US 08/973,462
PRIOR FILING DATE: 1998-02-06
PRIOR APPLICATION NUMBER: PCT/FR96/00894
PRIOR APPLICATION NUMBER: PCT/FR96/00894
PRIOR APPLICATION NUMBER: PCT/FR96/00894
                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/11/196,400
CURRENT FILING DATE: 2005-08-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: DAUBERŠIES, PIERRE
TITLE OF INVENTION: MALARIAL PRE-ERYTHROCYTIC STAGE POLYPEPTIDE MOLECULES
FILE REFERENCE: 200773USODIV
                                                                                                                          LENGTH: 20
TYPE; PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: FR 95/07007 PRIOR FILING DATE: 1995-06-13
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4; Conservative
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                    21.0%;
4; Mismatches
                 Score 26; DB 7;
Pred. No. 2.4e+02;
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Pred. No. 2.2e+02
2; Mismatches
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3; Indels
                                        Length 20;
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Gaps
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; TYPE: PRT ; ORGANISM: Homo sapiens US-11-054-515-2823
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US-11-054-515-2823
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; TYPE: PRT
; ORGANISM: Chlamydia pneumoniae
US-11-041-893-54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR PELICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-11-16
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Publication No. US20060002941A1
GENERAL INFORMATION:
APPLICANT: Mahairas, Gregory G.
TITLE OF INVENTION: COMPOSITIONS COMPRISING IMMUNE RESPONSE
TITLE OF INVENTION: ALTERING AGENTS AND METHODS OF USE
                                                                      Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2823
LENGTH: 12
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 54
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Best Local Similarity 62.5%;
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PFS23P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
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CURRENT EPLLICATION NUMBER: US/11/041,893
CURRENT FILING DATE: 2005-01-24
PRIOR APPLICATION NUMBER: US 60/616,855
PRIOR PILING DATE: 2004-10-06
PRIOR PILING DATE: 2004-01-23
PRIOR FILING DATE: 2004-01-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/543,296 PRIOR FILING DATE: 2004-02-11
                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
                                                                                                                                                                                                                           FILING DATE: 2001-03-21
APPLICATION NUMBER: 60/276,248
FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/293,499 FILING DATE: 2001-05-25 APPLICATION NUMBER: 60/277,379
                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 2001-06-15
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 09/880,748
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FILE REFERENCE: MER-133
CURRENT APPLICATION NUMBER: US/10/511,559
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR APPLICATION NUMBER: EP 03006554.4
PRIOR APPLICATION NUMBER: EP 03006554.4
PRIOR APPLICATION NUMBER: EP 03006554.4
PRIOR FILING DATE: 2003-03-24
NUMBER OF SEQ ID NOS: 1147
SOFTWARE: FAStSEQ for Windows Version 4.0
SEQ ID NO 256
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Best Local Similarity
Thehes 5; Conserve
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Best Local Similarity
Matches 5; Conserve
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; OTHER INFORMATION: Potential Epitope of human Factor VIII
US-10-511-559-256
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Search completed: January 26, 2006, 08:39:06
Job time : 4.55172 secs
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APPLICANT: JONES, Tim
APPLICANT: BAKER, Matthew
APPLICANT: CARR, Francis, J.
TITLE OF INVENTION: MODIFIED FACTOR VIII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 256, Ap
Publication No.
                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                    LENGTH: 13
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T-cell receptor al unidentified 27.2K stylar glycoprotei stylar glycoprotei actin - sea urchin Ig heavy chain DJ T-cell receptor be glutaminase - Alca capsid protein VP5 lipopolysaccharide protein kinase C i lens intrinsic mem cytochrome c3 - De high conductance c calmodulin, vasoaci type II site-speci acetylcholine rece equinatoxin 1A - s T-cell antigen rec wd-repeat protein cAMP-regulated pho creatine kinase (E comosain (EC 3.4.2 Ig H chain V-D-J r very-high-density T cell receptor be 13K protein 5306 { recombination prot ribulose-bisphosph cytochrome b559 co ribosomal protein clotting protein -mosact - sea urchi 14-3-3 protein hom amine oxidase (cop spectrin alpha cha celf protein - Esc polistes mastopara S-allele-associate S-allele-associate S-allele-associate GTP-binding protei T-cell antigen rec monodehydroascorba ananain (EC 3.4.22 hemocyanin chain I stylar glycoprotei protein 425 - Cali nitrogenase cofact calreticulin, protein-tyrosine k substance P - Atla calreticulin, hepa acetolactate synth protein kinase C i glutathione dehydr microbial serine I T cell receptor V 4-hydroxybenzoate major fat-globule neurotoxin-associa Ig heavy chain DJ amine oxidase (cop Ig heavy chain DJ urinary tract ston | pho |e (E |.4.2

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              R;Li, N.; Warren, P.V.; Golbeck, J.H.; Frank, G.; Zuber, H.; Bryant, D.A. Biochim. Biophys. Acta 1059, 215-225, 1991
A;Title: Polypeptide composition of the Photosystem I complex and the Photosystem I core A;Reference number: A58817; MUID:91355213; PMID:1853017
A;Accession: C56617
                                                                                                                                  photosystem I protein PsaL - Synechococcus sp. (PCC 6301) (fragment)
C;Species: Synechococcus sp.
C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 18-Aug-1995
C;Accession: C56817
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R;Crickmore, N.; Wheeler, V.C.; Ellar, D.J.

Mol. Gen. Genet. 242, 365-368, 1994

Mol. Gen. Genet. 242, 365-368, 1994

A;Title: Use of an operon fusion to induce expression and crystallisation of a Bacillus

A;Reference number: $42282; MUID:94150472; PMID:7906381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic Acids Res. 28, 862-867, 2000
A;Title: DNA sequence analysis of the photosynthesis region A;Reference number: Z25222; MUID:20115911; PMID:10648776
A;Accession: T50757
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C;Keywords: delta-
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A:Cross-rimental source: subsp. galleriae 916
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A;Molecule type: DNA
A;Residues: 1-20 <CHO>
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A;Experimental source: strain 2.4.1
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gene: cryIIB
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Pred. No. 3.2e+02;
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPARC:UPI0000115FBE; EMBL:X60865; NID:g53624; PIDN:CAA43255.1; A;Experimental source: T lymphocyte C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: T cell receptor genes in a series of class I major histocompatibility complex-r allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0771
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                                                                                                                              A;Genome: cr
C;Keywords:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 05-Nov-1999
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A; Residues: 1-21 < LI1>
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A;Experimental source: cv. 150
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A; Residues: 1-17 < KW2>
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A; Residues: 1-17 < KW1>
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A; Accession: JQ2320
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A; Residues: 1-12 < CAS>
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                                                                                                                                                      ;Genome: chloroplast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                             Query Match
Best Local
                                                                                                                                                                           Genetics
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                                                       Similarity 4; Conserv
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Pred. No. 1.2e+03;
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Pred. No. 1e+03;
2; Mismatches
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PID

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R;Kawagoe, Y.; Kikuta, Y.
Theor. Appl. Genet. 81, 13-;
A;Title: Chloroplast DNA evo
A;Reference number: JQ2306
A;Accession: JQ2310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: protein
A;Rosidues: 1-7,'X',9-19 <JAN>
A;Rossidues: 1-7,'X',9-19 <JAN>
A;Rosidues: 1-7,'X',9-19 <JAN>
A;Rosidues: 1-7,'X',9-19 <JAN>
A;Rosidues: UNIRAOT:Q9TRK6; UNIPARC:UPI000008701B; PIDN:AAB24426.1;
A;Experimental source: kidney
A;Note: sequence extracted from NCBI backbone (NCBIP:119897)
A;Note: we show one of the unidentified residues as tyrosine forming the top C;Superfamily: amiloride-binding protein
C;Keywords: oxidoreductase; quinoprotein; topaquinone
C;Keywords: oxidoreductase; quinoprotein; topaquinone
F;8/Modified site: topaquinone (Tyr) #status experimental
hypothetical protein K (pufQ 3' region) - Rhodobacter sphaeroides C;Species: Rhodobacter sphaeroides C;Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change 03-May-1994 C;Accession: $18582; $32885 R;Hunter, C.N.; McGlynn, P.; Ashby, M.K.; Burgess, J.G.; Olsen, J.D. Mol. Microbiol. 5, 2649-2661, 1991
A;Title: DNA sequencing and complementation/deletion analysis of the bcha-pu A;Reference number: $18580; MUID:92140030; PMID:1779756
A;Accession: $18582
                                                                                                                                                                                                            RESULT 8
S18582
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A;Title: Identification of topaquinone and
A;Reference number: A44239; MUID:93090748;
A;Accession: A44239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amine oxidase (copper-containing) (EC 1.4.3.6), kidney - pig (fragment) C;Species: Sus scrofa domestica (domestic pig) C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004 C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
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A; Residues: 1-17 < KAW>
A; Residues: 1-17 < KAW>
CTOOS- references: UNIPROT: 07M2E9; UNIPARC: UPI000017B098
A; Experimental source: strain Toko
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;Janes, S.M.; Palcic, M.M.; Scaman, C.H.; Smith, A.J.; Brown, D.E.; Dooley, D.M.;
;Ochemistry 31, 12147-12154, 1992
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Pred. No. 1.7e
1; Mismatches
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PMID:1457410
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1.7e+03;
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RESULT 11
B57789
C;Species: Homo sapiens (man)
C;Date: 23-Feb-1996 #sequence_revision 23-Feb-C;Accession: B57789
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A;Residues: 1-20 <HUN>
A;Cross-references: UNIPARC:UPI000017AB8B; EMBL:X68795
                                                                                                                                                                                                                                                                                                                                                                                J. Protein Chem. 10, 415-425, 1991
A;Title: Clostridium botulinum types A, B,
A;Reference number: A44644; MUID:92143938;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             neurotoxin-associated protein type B Hn+ 35K chain, band 3a - Clostridium botulinum C;Species: Clostridium botulinum C;Species: 07-0ct-1994 #sequence_revision 07-0ct-1994 #text_change 09-Jul-2004
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A;Cross-references: Unparc:UPI0000115262;
A;Note: this sequence was determined from t
C;Keywords: T-cell receptor
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C;Species: Mus musculus (house mouse)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change
                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:Q9R5N6; UA;Note: sequence extracted from NCBI
                                                                                                                                                                                                                                                                                                      A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-10 <SOM>
                                                                                                                                                                                                                                                                                                                                                    A;Contents: type B
A;Accession: F44644
                                                                                                                                                                                                                                                                                                                                                                                                                               C;Accession: F44644
R;Somers, E.; DasGupta, B.R.
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                                                                                                                                                                                                                                                       ;Note: sequence extracted ;Keywords: hemagglutinin
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Pred. No. 1.4e+03;
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Pred. No. 2e+03;
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RESULT 12
C36730
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J. Bacteriol. 172, 5477-5481, 1990
A;Title: Nuclectide sequence of the gene encoding the repressor A;Reference number: A36730; MUID:90368611; PMID:2203754
A;Accession: C36730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Klebsiella pneumoniae
C;Date: 19-Apr-1991 #sequence_revision 19-Apr-1991 #text_change 08-Oct-1999
C;Accession: C36730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Description: The proteins A;Reference number: A57789 A;Accession: B57789
                                                                                                                                                                                                                                                                                                                                             glycoprotein component 16/major fat-globule membrane protein/MFG-E8 homolog C;Species: Bos primigenius taurus (cattle) C;Decies: Bos primigenius taurus (cattle) C;Date: 19-Nov-1993 #sequence_revision 18-Nov-1994 #text_change 07-Feb-1997 C;Date: 19-Nov-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-9 <SCH>
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A,Molecule type: protein
A,Residues: 1-18 -GHN
A,Cross-references: UNIPARC:UPI000017C28A
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A;Title: The major fat-globule
II-like sequences.
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                                                                                                                                                              A; Experimental source: milk
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                                                                                                                                                                                                                              A;Accession: F48394
A;Status: preliminary
                                                                                                                                                                                                                                                              A;Reference number: A48394; MUID:93250576; PMID:8485470
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                                                                                                                                ;Keywords:
                                                                                                                                                                              ;Cross-references: UNIPARC:UPI000014C319
                                                                              Query Match
Best Local
                                                                                                                                                 Note: sequence extracted from NCBI backbone (NCBIP:131455)
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Best Local
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                                IOSVLNNGATR 23
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Protein Sequence Database,
he proteins of gallbladder
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29, 545-554, 1993
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36.4%;
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Pred. No. 2.
                                                               Score 23; DB 2;
Pred. No. 2.8e+03;
3; Mismatches 4
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C;Accession: S00494
R;Makino, N.; Kimura, S.
Eur. J. Biochem. 173, 423-430, 1988
A;Title: Subunits of Panulirus japonicus hemocyanin. 1.
A;Reference number: S00492; MUID:88196131; PMID:3360019
A;Accession: S00494
                                                                                                                                                                                                                                                                                           hemocyanin chain II - Japanese spiny lobster (fragment)
C;Species: Panulirus japonicus (Japanese spiny lobster)
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change
                                                                                                                                                                                                                                                                                                                                                    RESULT
S00494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene 129, 41-49, 1993
A;Title: Sequence analysis of the left end of the Bacillus subtilis bacteriophage SPP1
A;Reference number: JN0729; MUID:93328123; PMID:8335259
A;Accession: JN0733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: phage SPP1
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 17-Mar-1999
C;Accession: JN0730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical 1.7K protein - N;Alternate names: hypotheti C;Species: phage SPP1
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A;Title: D-arabinose dehydrogenase and biosynthesis of erythroascorbic acid A;Reference number: S72217; MUID:96439039; PMID:8841374
A;Accession: S72217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         D-arabinose 1-dehydrogenase [NAD(P)] (EC 1.1.1.117) - yeast (CarC)Species: Candida albicans
C;Date: 29-Jan-1998 #sequence_revision 06-Feb-1998 #text_change
C;Accession: S72217
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S72217
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A; Residues: 1-15 < CHA>
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A; Residues: 1-14 < KIM>
ş
                                                                                                                                    A;Molecule type: protein
A;Residues: 1-20 <MAK>
A;Cross_references: UNIPROT:P82312; UNIPARC:UPI000012C37C
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                                                                                                       C; Keywords:
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Best Local &
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Best Local
                                                                                                                        Superfamily: hemocyanin;
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5; Conserv
                                                                                                     copper; hemolymph; hexamer; oxygen carrier
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MAQNYKYRQQFIQSVLN 18
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hypothetical protein 42.1
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                                  Score 23; DB
Pred. No. 4e+0
6; Mismatches
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Pred. No.
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l; Mismatches
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R;Wallace, R.A.; Hoch, K.L.; Carnevali, O.
J. Mol. Biol. 213, 407-409, 1990
A;Title: Placement of small lipovitellin subunits within the vitellogenin precursor in A;Reference number: $10624, MUID:90278951; PMID:2352275
A;Accession: $10624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Experimental source: human bile and C; Superfamily: translation elongation C; Keywords: GTP binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Binette, J.P.; Binette, M.B.
submitted to the Protein Sequence Database, October 1996
A;Description: The proteins of kidney and gallbladder stones.
A;Reference number; A58501
A;Accession: D58503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C'Species: Becherichia coli
C'Species: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_char
C'Accession: PS0028
C'Accession: PS0028
J. Bacteriol. 171, 2728-2734, 1989
A.Title: DNA sequence analysis, gene product identification, A;Reference number: PS0027; MUID:89213963; PMID:2651416
A.Accession: PS0028
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S10624
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D58503
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A;Residues: 1-20 <MAL>
A;Cross-references: UNIPARC:UPI000017AA3D
C;Genetics:
                A; Molecule type: protein A; Residues: 1-12 <WAL>
                                                                                                                                       C;Accession: S10624
R;Wallace, R.A.; Ho
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C;Species: Escherichia coli
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 18-Jun-1993
                                                                                                                                                                lipovitellin - African clawed frog
7.Species: Xenopus laevis (African clawed frog)
7.Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references: UNIPROT:07M151; UNIPARC:UPI000017827D; Experimental source: human bile and stones
Cross-references: UNIPARC:UPI000017BF94
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Best Local
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Pred. No. 2.8e
0; Mismatches
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Pred. No. 4e+0
1; Mismatches
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J. 2.8e+05;
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4e+03;
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R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L. J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibe allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
                                                                                              T-cell receptor beta chain (M2) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 05-Nov-1999
C;Accession: PH0768
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PH0768
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Reference number: A; Accession: PS0454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            submitted to JIPID,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Tsugita,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Accession:
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C;Species: Oryza sativa (rice)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 23-Mar-1995
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PS0454
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jull-1992 #sequence_revision 17-Jul-1992 #text_change 05-Nov-1999
C;Accession: PH0746
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A, Title: T cell receptor genes in a series of class I major histocompatibility complex-
allelic exclusion and antigen-specific repertoire.
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J. Exp. Med. 174, 1371-1383, 1991
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Best Local S
Matches 3
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Similarity 40.0%;
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PS0206
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Pred. No. 4.9e+03;
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Pred. No. 3.4e+03;
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RESULT 23
S17461
S17461
Claredoxin B - Azotobacter chroococcum (fragment)
Clare: 19-Mar-1997 #sequence_revision 30-Jan-1998 #text_change 09-Jul-2004
Clate: 19-Mar-1997 #sequence_revision 30-Jan-1998 #text_change 09-Jul-2004
Claccession: S17461
R;Bagby, S.; Barker, P.D.; Hill, H.A.O.; Sanghera, G.S.; Dunbar, B.; Ashby, G.A.; Eady, Biochem. J. 277, 313-319, 1991
A;Title: Direct electrochemistry of two genetically distinct flavodoxins isolated from A;Reference number: S16929; MUID:91315397; PMID:1859358
A;Recession: S17461
A;Molecule type: protein
A;Residues: 1-20 SBAG-
                       RESULT 25
D42842
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A;Residues: 1-18 <CAS>
A;Residues: 1-18 <CAS>
A;Cross-references: UNIPARC:UPI0000115FBB; EMBL:X60862; NID:g52972; PIDN:CAA43252.1;
A;Experimental source: T lymphocyte
                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:P07475; UNIPARC:UPI00001763AB A;Note: proteinase inhibitor DE-3 has an estimated molecular weight C;Superfamily: plant Kunitz-type proteinase inhibitor C;Keywords: proteinase inhibitor
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JA0142
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antifungal 2S storage albumin large chain - radish (fragment)
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A;Title: Purification and properties of proteinase inhibitors from Erythrina corallodence A;Reference number: JA0142
A;Recession: JA0142
A;Recession: JA019
A;Residues: 1-20 <JOU>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Joubert, F.J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       proteinase inhibitor DE-3 - coral tree (fragment)
C;Species: Erythrina corallodendron (coral tree)
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 09-Jul-2004
C;Accession: JA0142
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C;Superfamily: flavodoxin; flavodoxin; flavodoxin; C;Kuperfamily: flavodoxin; flavoprotein; FYN
C;Keywords: electron transfer; flavoprotein; FYN
F;5-20/Domain: flavodoxin homology (fragment) <FLX>
F;9-14/Region: FYN-phosphate binding #status predicted
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Best Local
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Best Local :
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Best Local 9
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Local Similarity 42.9%;
les 3; Conservative
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57.1%;
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57.1%;
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Pred. No. 5.8e+03;
1; Mismatches 2;
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Pred. No. 5.2e+03;
2; Mismatches 2;
                                                                                                                                                                                                   1; Mismatches
                                                                                                                                                                                                                          Score 22; DB 2;
Pred. No. 5.8e+03;
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A;Note: sequence extracted from NCBI backbone (NCBIP:109925) C;Superfamily: Alpha amylase inhibitor
                                                                                                                                                                                                                                                                                                                                                              C;Species: Raphanus sativus (radish)
C;Date: 27-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 31-Dec-2004
C;Accession: D42842
C;Accession: D42842
R;Terras, F.R.; Schoofs, H.M.; De Bolle, M.F.; Van Leuven, F.; Rees, S.B.; Vanderleyden, J. Biol. Chem. 267, 15301-15309, 1992
A;Title: Analysis of two novel classes of plant antifungal proteins from radish (Raphanus A;Reference number: A42842; MUID:92348373; PMID:1639777
밁
                                                                                                                                                                                                                                                                        A; Molecule type: protein
A; Residues: 1-20 < TER>
                                                                                                                                                                                                                                                                                                                     A; Accession: D42842
A; Status: preliminary
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                                                                                                                                                                                                                            A; Experimental source: seed
                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q9S8Y7; UNIPARC:UPI00000A93E6
                                                                                        Matches
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Pred. No. 5.8e+03;
2; Mismatches 9;
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Search completed: January 26, 2006, 08:05:01 Job time : 13 secs

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Q7D25 FREDI
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Best Local
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                           conditions (By similarity).

-!- SUBUNIT: Oligomer of 14 subunits composed of to 15 subunits (By similarity).

-!- INDUCTION: By heat shock.
-!- SIMILARITY: BELONGS TO THE CHAPERONIN (HSP60) |
GO; GO:0005524; F:ATP binding; IEA.
GO; GO:00051082; F:unfolded protein binding; IEA.
GO; GO:0006457; P:protein folding; IEA.
GO; GO:0006986; P:response to unfolded protein; IE.
GO; GO:0006986; P:response to unfolded protein; IE.
InterPro; IPR001844; Chaprnin Cpn60.
PROSITE; PS00296; CHAPERONINS CPN60; PARTIAL.
ATP-binding; Chaperone; Heat shock.
NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Browne N., Dowds B.C.A.;
Submitted (JUL-2001) to Swiss-Prot.
Submitted (JUL-2001) to Swiss-Prot.
-i- PUNCTION: Prevents misfolding and promotes the refolding and
proper assembly of unfolded polypeptides generated under stress
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2001 (TrEMBLrel. 18, 01-OCT-2001 (TrEMBLrel. 18, 01-OCT-2003 (TrEMBLrel. 25,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
Hypothetical protein (Fragment).
ORFNames=PB402455.00.0;
ORFNames=PB402455.00.0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=NCIMB 11796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROTEIN SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=1396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacillus cereus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacillus cereus.
Bacteria; Firmicutes; Bacillales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P83074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P83074_BACCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preliminary data.
EMBL; CAAI01004758; CAI02160.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=5821;
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karyota; Alveolata; Apicomplexa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  kDa chaperonin (Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          w
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SSGYVFINLFIQKILN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AQNYKYRQQFIQSVLN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AND INDUCTION
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37.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cpn60) (groEL protein)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Created)
Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AF1F9ECC90CFE6FA CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacillaceae; Bacillus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genomic_DNA.
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RESULT 4
PUFK_RHOSH
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Q9S8A2_
RREFERENCE REPORTED TO THE PROPERTY OF THE PRO
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Best Local :
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                           Q53121; O08033;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (1)
PROTEIN SEQUENCE.
MEDLINE=96195148; PubMed=8660304;
Tatham A.S., Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; aster Campanilids; Asterales; Asteraceae; Asteroideae; Heliantheae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2000 (TrEMBLrel. 14,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
STRAIN=ATCC 17023 / 2.4.1 / NCIB
MEDLINE=96349111; PubMed=8760918;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biochem. J. SEQUENCE :
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Q9S8A2;
                                                                                                                                                      "Translational control of
                                                                                                                                                                                                                                                                                      NCBI_TaxID=1063;
                                                                                                                                                                                                                                                                                                                                                                                           Transcriptional regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sunflower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Millichip M.,
Stobart A.K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Helianthus annuus (Common
                  "R.sphaeroides genes
Submitted (MAR-1999)
                                                     McGlynn P.;
                                                                        NUCLEOTIDE SEQUENCE.
STRAIN=ATCC 17023 / 2.4.1 / NCIB 8253 / DSM
                                                                                                                                                                                                                                                                                                               Rhodobacteraceae; Rhodobacter.
                                                                                                                                                                                                                                                                                                                                   Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                      Rhodobacter sphaeroides (Rhodopseudomonas sphaeroides)
                                                                                                                                                                                                                                                                                                                                                                        Name=putK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PUFK_RHOSH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Purification and characterization of oil-bodies (oleosomes)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=4232;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          boundary proteins (oleosins) from lower (Helianthus annuus L.)."; hem. J. 314:333-337(1996).
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                                                                                                                                   142:2057-2069(1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1619 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2307
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46.7%;
              bchC, bchX, bchY, bchZ and pufQ.";
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                                                                                                                                                                                                                                                                                                                                   Alphaproteobacteria;
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                                                                                                                                                                       puf operon
                                                                                                                                                                                                                                                                                                                                                                               protein pufK.
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Last annotation update)
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Pred. No. 1
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Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDC28A062F7F8704 CRC64;
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                                                                                                                                                                                                                                  8253
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Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,

Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,

Berriman M., Florens L., Janssen C.S., Pain A., Christophides G.K.,

James K., Rutherford K., Harris B., Harris D., Churcher C.,

James K., Ormond D., Doggett J., Trueman H.E., Mendoza J.,

Place J. J., Parrell B., Turner C.M.R., Waters A.F., Sinden R.S.,

"A comprehensive survey of the Plasmodium life cycle by genomic,

transcriptomic, and proteomic analyses.",

Science 307:82-86(2005).
                                                                                                                                                                                                                                                                                                                                                                            13-SEP-2005 (TrEMBLrel. 31, Creat
13-SEP-2005 (TrEMBLrel. 31, Last
13-SEP-2005 (TrEMBLrel. 31, Last
Hypothetical protein (Fragment).
ORFNames=PB105963,00.0;
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Eukaryota; Alveolata;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Choudhary M., Kaplan S.;
"INA sequence analysis of the photosynthesis region of Rhodobacter sphaeroides 2.4.1.";
Nucleic Acids Res. 28:862-867(2000).
-!- FUNCTION: Involved in the transcriptional regulation of pufB.
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STRAIN-ATCC 17023 / 2.4.1 / NCIB 8253 / DSM 158;
MEDLINE-20115911; Pubmed-10648776; DOI=10.1093/nar/28.4.862;
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01-MAR-2004 (TrEMBLrel. 26, Last sequence up
01-MAR-2004 (TrEMBLrel. 26, Last annotation
Hypothetical protein (Fragment).
Name=PY06186;
                                                                                                                                                                                                                                                                                                                                                       01-MAR-2004
01-MAR-2004
01-MAR-2004
                                                          MEDLINE-22255706; PubMed=12368865; DOI=10.1038/nature01099; Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M. Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L., Selerson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M., Plorens L., Yates J.R. III, Raine J.D., Sinden R.E., Harris M.A., Cunningham D.A., Preiser P.R., Bergman L.W., Valdya A.B., van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R., Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.
"Genome sequence and comparative analysis of parasite Plasmodium yoelii yoelii."; Nature 419:512-519(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=22255706; PubMed=12368865; DOI=10.1038/nature01099; Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M., Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L., Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L., Peterson J.D., Pop M., Kosack D.S., Shunway M.F., Bidwell S.L., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Cho J.K., Quackenbush J., Sedegah M., Shoabib A., Cummings L.M., Florens L., Yates J.R. III, Raine J.D., Sinden R.E., Harris M.A., Cunningham D.A., Preiser P.R., Bergman L.W., Vaidya A.B., van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R., Valdya A.B., Van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R., Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.
                                                                                                                                                                                                                                                                                                                         Hypothetical Name=PY02751;
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Q7RKZ7;
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Eukaryota; Alveolata; Api
NCBI_TaxID=73239;
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Eukaryota; Alveolata; Api
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Q7RBF8;
Q1-MAR-2004 (TrEMBLrel. 2:
01-MAR-2004 (TrEMBLrel. 2:
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Q38371;
01-NOV-1996 (
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01-NOV-1996 (TrEMBLrel. 01, Last sequence
01-OCT-2003 (TrEMBLrel. 25, Last annotation coat protein, NCBI gi: 553004 (Fragment).
Bacteriophage MS2.
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EMBL; AABL01000761; EAA22236.1; -;
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Borrelia burgdorferi (Lym
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MEDLINE=73163785; PubMed=4512458;
Contreras R.R., Ysebaert M., Jou W.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical protein.
                                                 Stevenson B., Casjens S., Rosa P.;
"Evidence of past recombination events
Erp antigens of Borrelia burgdorferi.";
Microbiology 144:1869-1879(1998).
                                                                                                                                                                                                                                                                                               Bacteria; Spirochaete
Borrelia burgdorferi
NCBI_TaxID=139;
                                                                                                                                                                                                                                                                                                                                                                                                  Plasmid cp32-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein gene and the intercistronic region.";
Nature New Biol. 241:99-101(1973).
EMBL; M25.187; AAA32258.1; -; Genomic_RNA.
HSSP; P03612; IMSC.
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"Bacteriophage Ms2 RNA: nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Viruses, ssRNA positive-strand viruses, Levivirus; Levivirus subgroup I.
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                                                                                                                                                                        MEDLINE=98361033; PubMed=9695920;
                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                        Spirochaetes; Spirochaetales;
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Pred. No. 2.6e+03;
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
01-JUN-2000 (TrEMBLrel. 16, Last annotation update)
                                                                                                                                                                                                                           PROTEIN SEQUENCE.

MEDILNE=93133115; PubMed=1283000;

MEDILNE=93133115; PubMed=1283000;

Jeanteur D. Glatsu N., Pattus F., Buckley J.T.;

Jeanteur D., Glatsu N., Pattus F., Buckley J.T.;

"Purification of Aeromonas hydrophila major outer-membrane
"Purification of Aeromonas hydrophila"
"Purification o
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Q9R5K0;
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Q9R5E1;
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                                                                                                                                                                                          Mol. Microbiol. SEQUENCE 20 A
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Aeromonadaceae; Aeromonas
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Berriman M., Florens
James K., Rutherford
                                                                                                              Plasmodium berghei.
Eukaryota; Alveolata;
                                                                                                                                                   Hypothetical protein (Fragment). ORFNames=PB100192.00.0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Electrophoresis 20:781-789(1999).
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                                                      NUCLEOTIDE SEQUENCE.
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InterPro; IPR001404; Hsp90.
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DOI=10.1002/(SICT)1522-2683 [19990101)20:4/5<781::AID-ELPS781>3.0.CO;2-A;
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MEDINE-91312388; PubMed=1857378; DOI=10.1016/0166-6851(91)90152-V;

Menz B., Winter G., Ilg T., Lottspeich F., Overath P.;

"Purification and characterization of a membrane-bound acid
phosphatase of Leishmania mexicana.";

phosphatase of Leishmania mexicana.";
                  Hall N., Karras
Berriman M., Flo
                                                                                                                    Plasmodium chabaudi.
Eukaryota; Alveolata;
                                                                                                                                                         Hypothetical protein (Fragment). ORFNames=PC405777.00.0;
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Q4X617;
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13-SEP-2005
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01-MAY-2000
                                                          NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                        13-SEP-2005
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Bukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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rPro; IPR000560; HisAc_phsphtse.
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Churcher

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RESULT 17
Q7WAZB3_SOLTU PRELIMINARY;
ID Q7WZB3_SOLTU PRELIMINARY;
AC Q7WAZB3;
DT 01-MAR-2004 (TrEMBLrel. 26, L4
DT 01-MAR-2004 (TreMBLrel. 26,
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Bidwell S.L.,
Janse C.J., Ba
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Science 307.82-86 (2005).
-!- CAUTION: The sequence shown here is derived !
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13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
Hypothetical protein (Fragment).
ORFNames=PBL06774.00.0;
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Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium
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Science 307:82-86(2005).
-I- CAUTION: The sequence shown here is derived.
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13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation updat
13-SEP-2005 (TrEMBLrel. 31, Last annotation updat
Hypothetical protein (Fragment).
ORFNames=PC101630.00.0;
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NUCLEOTIDE SEQUENCE.

Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A., Berriman M., Florens L., Janssen C.S., Pain A., Christophid James K., Rutherford K., Harris B., Harris D., Churcher C., Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J. Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kaf Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden
                                                                                                                                                                                                                                                                                                                                                                             Q4Y571 PLACH PRELIMINARY;
Q4Y571;
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Vandekerckhove J., Ampe C.;
"Bukaryotic cytosolic chaperonin contains
seven related subunits.";
Proc. Natl Acad. Sci. U.S.A. 90:11975-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
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NUCLEOTIDE SEQUENCE.
Kikuta Y.;
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MEDLINE=94089752; PubMed=7903455;
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B608F6EBB5A8A2A6 CRC64;
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MEDILINE-21839745; PubMed=11682035; DOI=10.1016/S0168-8278(01)00167-2;

Kagaya M., Kaneko S., Ohno H., Inamura K., Kobayashi K.;

"Cloning and characterization of the 5'-flanking region of human cytokeratin 19 gene in human cholangiocarcinoma cell line.";

J. Hepatol. 35:504-511(2001).

EMBL; AB045973; BA8407700.1; -; Genomic_DNA.
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QSG554;
                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Mysticeti; Balaenopteridae; Balaenoptera.
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Science 307:82-86(2005).
-!- CAUTION: The sequence shown
EMBE/GenBank/DDBJ whole geno
                              NUCLEOTIDE SEQUENCE
                                                                                  NCBI_TaxID=9771;
                                                                                                                                                                                                    Balaenoptera musculus
                                                                                                                                                                                                                                         Name=POLB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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Q9BYF9;
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BALMU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GO:0005882; C:intermediate
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   C.A., May B.P.;
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66.7%;
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Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                    Created)
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 56BBD6970F8923EE CRC64;
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filament; IEA.
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Kawagoe Y., Kikuta ...

"Chloroplast DNA evolution in pocace."

Theor. Appl. Genet. 81:13-20(1991).

PIR, JQ2310, JQ2310.

PIR, JQ2310, JQ2310.
                                                                                                                                       Bukaryota; Viridiplantae; Strep
Spermatophyta; Magnoliophyta; e
lamiids; Solanales; Solanaceae;
                                                                             NUCLEOTIDE SEQUENCE.
Kawagoe Y., Kikuta Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in an African-American family.";
Hemoglobin 19:375-388(1995).
EMBL; S82767; AAD14420.1; -; Genomic_DNA.
HSSP; P02023; 1C7C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Global Population Structure of Blue Whales Balaenoptera musculus ssp Based on Nuclear Genetic Variation.";
Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; AY851160; AAW65802.1; -; Genomic_DNA.
NON_TER 1 1 16 16
                                                                                                                       NCBI_TaxID=4081;
                                                                                                                                                                                                   Lycopersicon esculentum
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Rahbar S., Nozari G., Forrest G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE
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Name=beta-globin;
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(TrEMBLrel. 13, Last sequence up
(TrEMBLrel. 26, Last annotation
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                                                                                                                                                                                                                                                                                             PRELIMINARY;
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Q9TWV6 APLCA PRELIMINARY; PRT; 18 AA.

ID Q9TWV6;
AC Q9TWV6;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 14, Last sequence update)
DT 01-MAY-2000 (TrEMBLrel. 14, Last annotation update)
DT 01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
DT 01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
DT 01-JUN-2000 (TrEMBLrel. 16, Last annotation update)
DT 01-JUN-2000 (TrEMBLrel. 17, Last annotation update)
DT 01-JUN-2000 (TrEMBLrel. 18, Last annotation update)
DT 01-JUN-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
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DT 01-JUN-2000 (TrEMBLrel. 14, Last sequence update)
DT 01-JUN-2000 (TrEMBLrel. 14, Last sequence update)
DT 01-JUN-2000 (TrEMBLrel. 13, Last sequence).

CS Aplysic californica (California sea hare).

CC Aplysic california sea hare).

CC Aplysicidea; Aplysidae; Aplysia.

OC Aplysicidea; Aplysidae; Aplysia.

(1)
RP PRTEIN SEQUENCE.
RY MEDLINE=93057436; PubMed=1431843;
RY MISON M.P., Carrow G.M., Levitan I.B.;
RY "Modulation of growth of Aplysia neurons by an endogenous lectin.";
RL J. Neurobiol. 23:739-750(1992).
SQ SEQUENCE 18 AA; 2105 MW; 5652840B573057D8 CRC64;
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8Y3 LUPAR PRELIMINARY;
Q958Y3;
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Q1-MAY-2000 (Tremblrel. 13,
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GO; GO:0004067; F:asparaginase activity; IEA.

SEQUENCE 17 AA; 1703 MW; 9AEDD9691F7F0807 CRC64;
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ol-MAY-2000 (TrEMBLrel. 13, Last sequence update)
ol-JUN-2003 (TrEMBLrel. 24, Last annotation update)
L-asparaginase isoform A (EC 3.5.1.1) (Fragment).
Lupinus arboreus (Tree lupin).
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MEDLINE-92344803; PubMed=1368361; DOI=10.1016/0031-9422(92)83098-J;
LOUGH T.J., Chang K.S., Carne A., Monk B.C., Reynolds P.H.,
Farnden K.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 20.2%; Score 25; DB 2; Local Similarity 60.0%; Pred. No. 9.2e+03;
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7 Human B-d
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9 Human hep
1 Lassa vir
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 BLys; B lymphocyte stimulator; TNF superfamily; human; cytostatic;
                     Human BLyS binding scFv VH CDR3 SEQ ID 2216
                                             19-AUG-2002
                                                                 ABP46205;
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                                                                                                                                                                                                                                                    detect HPV-infection.
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                                                                                                                                                                                                                                                                                                                                                                        Diagnosis of human papilloma virus infection and PV-carrying tumours using synthetic peptide(s) to detect virus specific antigen-antibody
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                                                                                                                                                                                      10;
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                                                                                                                                           σ
                                                                                                                                                                                                Similarity
                                                                                                                                           EDNVMEYVVWT-HIYL 19
                                                                                                                                                               EXNVMXYVLYTMHYYL 17
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                                                                                                                                                                                                                                                                                                                                                                                                                             Dillner L,
                                                                                                                                                                                                                                AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25.4
25.4
                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                               immunoassay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90SE-00001705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90SE-00001705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cervical intraepithelial neoplasia; CIN; wart;
                                                                                      peptide;
                                            entry)
                                                                                                                                                                                               37.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16
16
                                                                                                                                                                                                                                                     See AAR15523-R15601
                                                                                                                                                                                                                                                                                                                                                                                                                              Cheng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADK40360
ADM12543
ADO38774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                       16
                                                                                                                                                                                     2
                                                                                                                                                                                     Score 44.5; DE Pred. No. 2.8; 2; Mismatches
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                                                                                                                                                                                                         Length 20;
                                                                                                                                                                                       Indels
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Adm12543
Ado38774
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3 Ii-key/hu
4 Human tyr
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RESULT 3
ADG97032
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Best Local Similarity
Matches 5; Conserv
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17-OCT-2000;
16-MAR-2001;
21-MAR-2001;
25-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic, immunosuppressive, immunostimulant, immunomodulatory, antirheumatic and antiAIDS activity and can be used in vaccines to inhibit the expression and activity of BLyS. The antibodies bind to BLyS and so may be used to detect and quantitate the presence of BLyS in biological samples and may be used in this way to diagnose disease associated with aberrant expression of BLyS. They may also be administered to treat diseases associated with aberrant BLyS expression and activity such as cancer, immune, and autoimmune disorders and diseases, e.g. systemic lupus erythematosus, rheumatoid arthritis, immunodeficiency (e.g. common variable immunodeficiency (CVID) and acquired immunodeficiency syndrome (AIDS). ABP4399-ABP47228 represent
antibody; B lymphocyte stimulator; BLyS; tumour necrosis factor;
B cell proliferation; differentiation; scFv; myasthenia gravis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel antibodies that immunospecifically bind to B Lymphocyte Stimulator (BLyS) polypeptides. BLyS is a member of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunosuppressive; immunostimulant; immunomodulatory; antirheumatic; antiAIDS; vaccine; cancer; immune; autoimmune disorder; immunodeficiency; systemic lupus erythematosus; rheumatoid arthritis; CVID; AIDS; common variable immunodeficiency; acquired immunodeficiency syndrome.
                                                                              11-MAR-2004
                                                                                                                                                                                                                                                                                                                                          Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour necrosis factor (TNF) super family and induces B cell proliferation and differentiation. The antibodies of the inv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                scFV VHCDR3
                                                                                                                                         ADG97032
                                                                                                                                                                                                                                                                                                                                                                                       acquired immunodeficiency syndrome (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antibodies against B Lymphocyte Stimulating polypeptides, useful for diagnosis and treatment of cancers and immune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUN-2001; 2001WO-US019110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour necrosis factor; B cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                          the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200202641-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HUMAN GENOME SCI INC
CAMBRIDGE ANTIBODY TO
                                                                                                                                         standard;
                                                                                                                                                                                                                                                TAXHWLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 2953; 3148pp; English.
                                                                                                                                                                                                                     TAAHTAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barash SC,
                                              peptide that immunospecifically binds BLyS SeqID 2216
                                                                                                                                                                                                                                                                                                                                            3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0240816P.
2001US-0276248P.
2001US-0277379P.
2001US-0293499P.
                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0212210P
                                                                             (first entry)
                                                                                                                                        peptide; 16
                                                                                                                                                                                                                                                                                             29.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Choi GH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TECHNOLOGY.
                                                                                                                                                                                                                                                                                             Score 35;
Pred. No.
                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vaughan T,
                                                                                                                                                                                                                                                                                                                                                                                          antibodies described in the method of
                                                                                                                                                                                                                                                                                               DB
84;
                                                                                                                                                                                                                                                                                                             <u>ت</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hilbert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B cell differentiation;
                                                                                                                                                                                                                                                                               1.
                                                                                                                                                                                                                                                                                                            Length 16;
                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention have
                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BLyS
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RESULT 4
AAE30892
ID AAE3
XX AAE3
XX AAE3
XX AAE3
XX AAE3
XX AAE3
XX DT 24-F
DT 24-F
DT 24-F
DT 25-F
D
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to novel antibodies that immunospecifically bind CC to B lymphocyte stimulator (BLyS). The BLyS gene has been mapped to CC chromosome 13934 and encodes a protein that is a member of the tumour necrosis factor superfamily and induces both in vivo and in vitro B cell CC proliferation and differentiation. Specifically, it refers to single CC chain antibody molecules (serve) derived, preferably, from the variable CC chain antibody molecules (serve) derived, preferably, from the variable CC resent invention refers to the use of such antibodies in various methods for the detection, diagnosis and prognosis of diseases related to the CC present invention refers to the use of such antibodies in various methods CC for the detection, diagnosis and prognosis of diseases related to the aberrant expression or inappropriate function of BLyS or its receptor. As concluding myasthenia gravis and multiple sclerosis, inflammatory compositions are useful for identifying immune disorders consolitions and rheumacoid arthritis, infectious diseases such CC disorders e.g. asthma and rheumacoid arthritis, infectious diseases such CC lymphoma. Accordingly, they can be described as exhibiting various activities such as antirheumatic, antiarthritic, neuroprotective, antiasthmatic, antiasthmatic, neuroprotective, antinflammatory, antiasthmatic, antiablegic and cytostatic. This coptide sequence is a single chain antibody variable heavy CDR3 peptide that immunospecifically binds BLyS of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
    Unidentified
                                                                p53 polypeptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     multiple sclerosis; asthma; rheumatoid arthritis; AIDS; leukaemia; carcinoma; lymphoma; antirheumatic; antiarthritic; neuroprotective; antiinflammatory; antiasthmatic; antiallergic; cytostatic.
                                            cancer; ischaemia;
                                                                                                         MDM2-binding
                                                                                                                                                     24-FEB-2003
                                                                                                                                                                                                AAE30892;
                                                                                                                                                                                                                                      AAE30892 standard; peptide; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel antibody that immunospecifically binds to a B lymphocyte stimulator (BLys), useful for detecting and treating diseases or disorders e.g. rheumatoid arthritis, asthma and leukemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-NOV-2001;
19-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                          1
                                                                                                                                                                                                                                                                                                                                                 9 YVLHYYL 15
                                                                                                                                                                                                                                                                                                                                                                                                                                    5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                          YTMHYYL 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Α
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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2001US-0340817P.
                                                                                                    motif #7, identified by phage-peptide display.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002WO-US036496
                                                                                                                                                     (first
                                                                p300 polypeptide; cell cycle;
                                            cytostatic; vasotropic
                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                      29.7%;
71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Choi GH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    394pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 35; DB Pred. No. 84; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vaughan TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English
                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7;
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                                                            cell death; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                D,
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ABP46248
  밁
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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16-JUN-2000;
17-OCT-2000;
16-MAR-2001;
21-MAR-2001;
25-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                       immunosuppressive; immunostimulant; immunomodulatory; antitheumatic; antiAIDS; vaccine; cancer; immune; autoimmune disorder; immunodeficiency; systemic lupus erythematosus; rheumatoid arthritis; CVID; AIDS; common variable immunodeficiency; acquired immunodeficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a psptide for use in modulating the binding of p53 polypeptide to a p300 polypeptide. The new peptide is useful in modulating the binding of a p53 polypeptide to a p300 polypeptide. The peptide may be used to regulate the mammalian cell cycle or to induce or prevent cell death, for the treatment of cancer or ischaemia. The invention is useful in gene therapy. The present sequence is MDM2-binding motif, identified by phage-peptide display
  (HUMA-)
                                                                                                                                                                                                                                                                                       WO200202641-A1
                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            BLys; B lymphocyte stimulator; TNF superfamily; human; cytostatic; tumour necrosis factor; B cell proliferation; B cell differentiati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABP46248 standard; peptide; 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New peptide for modulating the binding polypeptide, useful for regulating the treatment of cancer or ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-FEB-2002;
                                                                                                                                                                                                15-JUN-2001; 2001WO-US019110
                                                                                                                                                                                                                                            10-JAN-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BLyS binding
HUMAN GENOME SCI
CAMBRIDGE ANTIBOI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 6; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dornan D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fig
                                                             ; 2000US-0212210P.
; 2000US-0240816P.
; 2001US-0276248P.
; 2001US-0277379P.
; 2001US-0293499P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002WO-GB000640
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  ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 scFv VH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28.8%;
50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDR3
  TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 34; DB
Pred. No. 88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID 2259
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6
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ADG97075
ID ADG9
XX ADG5
XX ADG5
XX ADG5
XX ADG5
XX ADG7
Il-N
XX ADG1
KW ADG1

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic, immunosuppressive, immunostimulant, immunomodulatory, antirheumatic and antiALIDS activity and can be used in vaccines to inhibit the expression and activity of BLyS. The antibodies bind to BLyS and so may be used to detect and quantitate the presence of BLyS in biological samples and may be used in this way to diagnose disease associated with aberrant expression of BLyS. They may also be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antibody; B lymphocyte stimulator; BLyS; tumour necrosis factor; B cell proliferation; differentiation; scFv; myasthenia gravis; multiple sclerosis; asthma; rheumatoid arthritis; AIDS; leukaemia; carcinoma; lymphoma; antirheumatic; antiarthritic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and activity such as cancer, Inmunic, and concommentation arthritis, diseases, e.g. systemic lupus erytheatnosus, rheumatoid arthritis, immunodeficiency (cyll) and immunodeficiency (e.g. common variable immunodeficiency (CVID) and acquired immunodeficiency syndrome (AIDS)). ABP43990-ABP47228 represent acquired immunodeficiency syndrome (AIDS)). ABP43990-ABP47228 represent acquired immunodeficiency syndrome (AIDS).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel antibodies that immunospecifically bind (B Lymphocyte Stimulator (BLyS) polypeptides. BLyS is a member of the tumour necrosis factor (TNF) super family and induces B cell proliferation and differentiation. The antibodies of the invention have
     Novel antibody that immunospecifically binds to a B lymphocyte stimulator
                                                              WPI; 2003-505530/47
                                                                                                                                                                                                                                 16-NOV-2001;
19-DEC-2001;
                                                                                                                                                                                                                                                                                                                    14-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003055979-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BCFV VHCDR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG97075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG97075 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     administered to treat diseases associated with aberrant BLyS expression and activity such as cancer, immune, and autoimmune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2; Page 2960;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-114799/15
                                                                                                                                                                                                                                                                                                                                                                             10-JUL-2003
                                                                                                                                                                            (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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MS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
5; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B against B Lymphocyte Stimulating
and treatment of cancers and immun
                                                                                                                   Barash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Barash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide that immunospecifically binds BLyS SeqID 2259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A,
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                                                                                                                                                                                                                                 2001US-0331469P.
2001US-0340817P.
                                                                                                                                                                                                                                                                                                                       2002WO-US036496
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                                                                                                                   SC,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3148pp;
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                                                                                                                   Choi GH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 34; DB
Pred. No. 1.2e
0; Mismatches
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                                                                                                                Vaughan TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 5;
1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ating polypeptides, useful for the immune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hilbert
                                                                                                                      Hilbert D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chromosome 13q34 and encodes a protein that is a member of the tumour necrosis factor superfamily and induces both in vivo and in viro B ce proliferation and differentiation. Specifically, it references chain antibody molecules factor.
                                                   Diagnosis of human papilloma virus infection and PV-carrying using synthetic peptide(s) to detect virus specific antigen-complexes by immunoassay.
                                                                                                                                                                                                                                                  28-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                            02-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR15580 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16
                                                                                                            WPI; 1991-369390/50
                                                                                                                                      Dillner J,
                                                                                                                                                                                               11-MAY-1990;
                                                                                                                                                                                                                        11-MAY-1990;
                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                    squamous cell
                                                                                                                                                                                                                                                                                                                                                      cervical
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR15580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BLys), useful for detecting rheumatoid arthritis, asthma
                                                                                                                                                                   (MEDS-) MEDSCAND
                                                                                                                                                                                                                                                                                                                                                                              Immunopeptide derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11 AAHWAA 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                    cancer; cervical intraepithelial neoplasia; CIN; wart;
cell carcinoma; ELISA; HPV 18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      YVMHYY 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Å
                                                                                                                                      Dillner L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                             90SE-00001705
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                                                                                                                                                                                                                         90SE-00001705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28.8%;
                                                                                                                                                                                                                                                                                                                                                                                from HPV18 E2 peptide
                                                                                                                                      Cheng
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and treating diseases and leukemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 34; DB 7;
Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  that immunospecifically bind
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 16;
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                                                                   -carrying tumours antigen-antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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Disclosure;

Page

39; 72pp; English

This is one

of a large number of peptides which have been synthesised on

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Best Local
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                                                                                                                                                                                                                                                                                               02-SEP-1997;
01-MAY-1998;
04-NOV-1998;
                                                                                                        Yacoby-Zeevi O,
Ayal-Hershkovitz
Friedmann Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the basis of the amino acid sequences for the E2, E4, E7, L1 or L2 proteins of HPV 1, 5, 6, 8, 11, 16, 18, 31 and 33. The selection of peptide sequences was based on the assumption that an immunoreactive region might be situated in the same relative region of a protein from different HPV types. The peptides were used in diagnostic immunoassays to detect HPV-infection. See AAR15523-R15601
                                                   Targeted drug delive useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-NOV-2004
                                                                                                                                                                                                         (SHLO/)
                                                                                                                                                                                                                               (PERE/)
(MIRO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Targeted drug delivery; inflammatory disorder; wound; scar; vasculopathy; autoimmune disorder; cancer; angiogenesis; metastatic disease; atherosclerosis; restenosis; aneurysm; solid cancer; non-solid cancer; haematopoietic malignancy; lymphocytic leukaemia; myelogenous leukaemia;
                                                                                     WPI; 2004-625084/60
                                                                                                                                                                                                                                                                          19-FEB-2003;
22-AUG-2003;
                                                                                                                                                                                                                                                                                                                                           28-NOV-2003; 2003US-00722502
                                                                                                                                                                                                                                                                                                                                                                                     US2004170631-A1
                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                              Hodgkin's disease;
human; heparanase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADR88216 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                    (YACO/)
                                                                                                                                                                                    FEIN/)
                                                                                                                                                                                               (AYAL/)
                                         cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           heparanase
                                                                                                                                                  VAN GELDER J M.
VLODAVSKY I.
FRIEDMANN Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -
                                                                                                                                                                                                                               PERETZ T
MIRON D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4
                                                                                                                                                                                               AYAL-HERSHKOVITZ
                                                                                                                                                                                                         PECKER I.
                                                                                                                                                                                                                      SHLOMI Y.
                                                                                                                                                                                                                                                    YACOBY-ZEEVI O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 6; Conserv
                                       drug delivery to a heparanase-expressing tissue of a patient, or treating heparanase-associated conditions such as inflammation r, comprises administering a drug and an anti-heparanase antibody
                                                                                                                                                                                     FEINSTEIN E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
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98US-00071739.
98US-00186200.
2003US-00368044.
2003US-00645659.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                    Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         epitope
                                                                                                                              Peretz T,
                                                                                                                                                                                                                                                                                                                                                                                                                               enzyme; epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                          multiple myeloma; haemangiosarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28.8%;
                                                                                                                    Feinstein
                                                                                                                                                                                               3
                                                                                                                 Miron D, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 34; DB
Pred. No. 1.6e
4; Mismatches
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                                                                                                                   Shlomi Y,
n Gelder JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.6e+02;
7;
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                                                                                                                 Pecker I;
, Vlodavsky I;
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The
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Claim 7;

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58pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
02-SEP-1997;
01-MAY-1998;
04-NOV-1998;
19-FEB-2003;
                                                                                                                                                                                                                                                                                                              Antibody; epitope; heparanase; pathological condition; angiogenesis; cell proliferation; cancerous condition; tumour cell invasion; metastatic disease; heparanase-related disorder; inflammatory disorder; wound; scar; vasculopathy; autoimmune condition; renal disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method of targeted drug delivery to a tissue of a patient, the tissue expressing heparanase. The method comprises providing a complex of a drug directly or indirectly linked to an antiheparanase antibody, and administering the complex to the patient. In the targeted drug delivery, the antibody comprises an antibody or its portion capable of specifically binding to at least one epitope of a heparanase protein. The composition and methods of the invention are useful for
                                                                                                                                                                                     US2004213789-A1
                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                        cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                     Functional peptide epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADT78183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADT78183 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      malignant melanoma of the conjunctiva, malignant melanoma of the uvea, retinoblastoma, carcinoma of the lacrimal gland, sarcoma of the orbit, brain, spinal cord, vascular system, haemangiosarcoma and Kaposi's
                                                                                                  22-AUG-2003; 2003US-00645659
                                                                                                                                            28-0CT-2004:.
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                                                                                                                                                                                                                               sapiens.
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                                                                                                                                                                                                                                                                      immunosuppressive;
                                                                                                                                                                                                                                                                                        antiinflammatory;
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                     98US-00071739
98US-00186200
                                                             97US-00922170
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72
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                                                                                                                                                                                                                                                                                                                                                                                                                     of human heparanase, pep10.
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Pred. No. 1.3e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                    vulnerary; antiarteriosclerotic;
; nephrotropic; antidiabetic; hum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       is human heparanase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     epitope
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RESULT 10
AEA42432
ID AEA42
XX AEA42
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                                                                                                                                                                                                                                                                                                                                                                                                                                       subject, (c) a method for monitoring the state of a heparanase-related condition in a subject, and (d) a pharmaceutical composition and comprising the isolated anti-heparanase antibody or antibody portion and a pharmaceutical carrier. The antibody, methods, and composition are useful for detecting, treating, preventing or monitoring a pathological condition, e.g. angiogenesis, cell proliferation, a cancerous condition (blood, breast, bladder, rectum, stomach, cervix, ovarian, colon, renal, or prostate cancer) minor cell proliferation, invasion of circulating tumour cells, or a metastatic disease, or a heparanase-related disorder or condition, e.g. an inflammatory disorder, wound, scar, vasculopathy (atheroselerosis, restenosis, or aneurym), autoimmune condition, or renal disease or disorder (diabetic nephrotic syndrome, or a renal cell carcinoma in a mammal. This sequence represents a functional peptide
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated antibody or antibody portion capable of specifically binding to or elicited by at least one epitope of a heparanase protein, where the heparanase protein is at least 60% homologous to any of the 6 sequences given as SEQ ID NOS: 1-5 or 11, and where at least one epitope comprises a sequence at least 70% homologous to any of the sequences given as SEQ ID NOS: 6-10. Also disclosed are (a) a hybridoma cell line comprising a cell line for producing the monoclonal antibody, (b) a method for detecting, treating or preventing a pathological condition or a heparanase-related disorder or condition in a
antibody; heparanase; antiinflammatory; vulnerary; in antiangiogenic; cytostatic; antiarteriosclerotic; vas inflammation; wound healing; scarring; vasculopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PERE/)
(MIRO/)
(SHLO/)
(PECK/)
(AYAL/)
                                                                          Human heparanase epitope peptide SEQ ID NO:10.
                                                                                                                 28-JUL-2005
                                                                                                                                                      AEA42432
                                                                                                                                                                                        AEA42432 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New neutralizing monoclonal anti-heparanase antibodies, useful for detecting, treating or preventing cancer, inflammatory or autoimmu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-774790/76.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yacoby-Zeevi O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (FEIN/)
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AYAL-HERSHKOVITZ |

FEINSTEIN E.

GELDER J M V.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of human heparanase.
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                                                                                                                                                                                                                                                                                      HYYLNGRTATR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , treating or preventing cancer, inflammatory or autoimmune wound, atherosclerosis, restenosis, or diabetic nephropathy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 10; 68pp; English.
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                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                               (first entry
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M, Feinstein
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sin E, Gelder JMV,
                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                            Score 33.5; DB 8;
Pred. No. 1.3e+02;
0; Mismatches 2;
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                   vasotropi
                                                                                                                                                                                                                                                                                                                                                                Indels
                                   immunosuppressive;
   autoimmune disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of specifically binding to an epitope of a heparanase protein. Also described: (1) a cell line (II) for producing a monoclonal antibody or its portion, comprising a cell line for producing (I); (2) a pharmaceutical composition comprising (I) and a carrier; and (3) an affinity medium (III) for binding human heparanase polypeptides, comprising (I) immobilized to a chemically inert, insoluble carrier. (I) useful for treating a subject suffering from a pathological condition, which involves administering (I) to the subject. (I) is useful for preventing and treating heparanase-related disorder or condition chosen
                                                                                                                                                                                                                                                                                                                                                                                                                                              from inflammatory disorder, wound, scar, vasculopathy, autoimmune condition, angiogenesis, cell proliferation, cancerous condition, tumor cell proliferation, invasion of circulating tumor cells and metastatic disease (I) is useful for detecting the presence of heparanase polypeptide in a sample. (I) is useful for detecting heparanase-related disease or condition in a subject such as vertebrate, preferably mammal e.g., human. The heparanase-related disease or condition further includes renal disease or disorder of condition further glomerulosclerosis, nephrotic syndrome, minimal change nephrotic syndrome and renal cell carcinoma. The present sequence represents a human fither and cell carcinoma. The present sequence represents a human fither and cell carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated antibody capable of specifically binding to epitope of heparanase protein, useful for preventing and treating heparanase-related disorder such as inflammatory disorder, scars, autoimmune conditions or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2005-173343/19
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Shlomi Y, Peleg Y,
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                                                                                                                                            ADU70841 standard;
                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  heparanase epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INSI-) INSIGHT BIOPHARMACEUTICALS LTD. (HADA-) HADASIT MEDICAL RES SERVICES &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          angiogenesis disorder; cancer; tumor; metastasis; epitope
                                                                                                                                                                                                                                                                     14 HYYLN-NGATR 23
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                                                                                                                                                                                                                                                                                                      Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  relates to an isolated antibody or its portion (I) capable
                                                                                                                                         peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide, which
                                                                                                                                                                                                                                     14
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Yacoby-Zeevi O,
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                                                                                                                                                                                                                                                                                                                         Score 33.5; DB 9; Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                    is used
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D, Ayal-Hershkovitz M,
                                                                                                                                                                                                                                                                                                                                            DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                    in the
                                                                                                                                                                                                                                                                                                                                        Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                 exemplification of the
                                                                                                                                                                                                                                                                                                          Indels
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10-FEB-2005

(first entry)

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Human heparanase peptide SEQ ID

NO:526

enzyme; heparinase; vaccine; human leukocyte antigen; HLA;

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RESULT 12
ADU71059
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Best Local S
Matches 8
WPI; 2005-014847/02
                                                                                                                                                                                                                                                                                                                                                                                                                    enzyme; heparinase; vaccine;
immunostimulant; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human heparanase peptide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADU71059;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel heparanase peptide that binds to a human leukocyte antigen (HLA) molecule, where the peptide is a nonapeptide, or its functional derivative. A peptide of the invention has immunostimulant and cytostatic activity, and is used in a vaccine. The heparinase peptide is useful for preparing a medicament which induces an immune response or for treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New heparanase nonapeptide that binds to a human leukocyte antigen (HLA) molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
                                            Schirrmacher V,
                                                                                                                                                                     19-MAY-2003; 2003EP-00011038
                                                                                                                                                                                                                                                                        24-NOV-2004.
                                                                                                                                                                                                                                                                                                                         EP1479764-A1
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                                                                                                                                                                                                                       19-MAY-2003; 2003EP-00011038
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nilarity 72.7%;
Conservative
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                                                                                          KREBSFORSCHUNGSZENTRUM
HEIDELBERG RUPRECHT-KAI
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                                              Beckhove
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                                                                                             RUPRECHT-KARLS
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Pred. No. 1.4e.
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                       human leukocyte antigen; HLA; immune disorder; metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ID NO:744
                                              Sommerfeldt
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les 2;
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ADU71060
ID ADU7
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AC ADU7
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Best Local !
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                            The invention relates to a novel heparanase peptide that binds to a human leukccyte antigen (HLA) molecule, where the peptide is a nonspeptide, or its functional derivative. A peptide of the invention has immunostimulant and cytostatic activity, and is used in a vaccine. The heparinase peptide is useful for preparing a medicament which induces an immune response or for treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
                                                                                                                                                                                          New heparanase nonapeptide that binds to a human leukocyte antigen (HLA) molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel heparanase peptide that binds to a human leukocyte antigen (HIA) molecule, where the peptide is a monapeptide, or its functional derivative. A peptide of the invention has immunostimulant and cytostatic activity, and is used in a vaccine. The heparinase peptide is useful for preparing a medicament which induces an immune response or treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
 Sequence 15 AA;
                                                                                                                                                              Disclosure; SEQ ID NO 745; 269pp; English.
                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                          Schirrmacher V,
                                                                                                                                                                                                                                                                                                                          (DEKR-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human heparanase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADU71060 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New heparanase nonapeptide that binds to a human leukocyte antigen (HLA) molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         24-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      enzyme; heparinase; vaccine;
immunostimulant; cytostatic;
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HEIDELBERG RUPRECHT-KARLS.
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                                                                                                                                                                                                                                                                                          Beckhove P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide
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72.7%;
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Pred. No. 1.4e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human leukocyte antigen; immune disorder; metastas
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                                                                                                                                                                                                                                                                                            Sommerfeldt
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Query Match

28

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Score 33.5;

DB 9;

Length

15;

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ADU7
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ADU71272
ID 72
ID 72
ID 72
ID 72
ID 72
ID 72
ID 74
ID 77
IO-FE
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IO-FE
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ID 78
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ADU70927
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Best Local S
Matches 8
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Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel heparanase peptide that binds to a human leukocyte antigen (HLA) molecule, where the peptide is a nonapeptide, or its functional derivative. A peptide of the invention has immunostimulant and cytostatic activity, and is used in a vaccine. The heparinase peptide is useful for preparing a medicament which induces an immune response or for treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New heparanase nonapeptide that binds to a human leukocyte antigen (HLA) molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
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                        Human heparanase peptide SEQ ID NO:612
                                                                    10-FEB-2005
                                                                                                            ADU70927;
                                                                                                                                                    ADU70927 standard; peptide; 15 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 957; 269pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human heparanase peptide
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UNIV HEIDELBERG RUPRECHT-KARLS.
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                                                                                                                                                                                                                                                                                                     HYYLN-NGATR 23
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                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; immune disorder; metastasis.
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0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                           Score 33.5; DB 9;
Pred. No. 1.4e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                     Length 15;
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  RESULT 16
ADY27047
ID ADY277
XX ADY27
XX ADY27
XX O5-MA
XX Hepax
XX Hepax
KW neuro
KW neuro
KW nepax
XX Synth
XX 12-AU
PR 11-AU
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Best Local S
Matches 8
                                                                  14-AUG-2003; 2003US-0494800P.
12-JAN-2004; 2004US-0535492P.
                                                                                                                                                                                                                                                                                               Heparanase; cancer; neoplasm; inflammation; cardiovascular disease; neurological disease; viral infection; infection; cytostatic; antiinflammatory; cardiovascular-gen.; neuroprotective; virucide; heparanase modulator; enzyme purification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel heparanase peptide that binds to a human leukocyte antigen (HLA) molecule, where the peptide is a nonapeptide, or its functional derivative. A peptide of the invention has immunostimulant and cytostatic activity, and is used in a vaccine. The haparinase peptide is useful for preparing a medicament which induces an immune response or for treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New heparanase nonapeptide that binds to a human leukocyte antigen (HLA) molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
                                                                                                                                                                                                                                                                                                                                                                                                             Heparanase inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADY27047;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP1479764-A1
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                                                                                                                                12-AUG-2004; 2004WO-IL000744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADY27047 standard; peptide; 15
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                          (INSI-) INSIGHT BIOPHARMACEUTICALS LTD
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UNIV HEIDELBERG RUPRECHT-KARLS.
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                                                                                                                                                                                                                                                                                                                                                                                                             peptide
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72.7%;
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Score 33.5; DB 9; Pred. No. 1.4e+02; 0; Mismatches 2

9;

Length 15; Indels

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Gaps

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Sommerfeldt

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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               heparanase, a protease substrate mimetic comprising a peptide representing a subset or all substrate residues or cleavage sites of human heparanase or an equivalent non-human heparanase, a method of producing active heparanase and a method of modulating an adhesion activity of heparanase. The composition and methods are useful for modulating heparanase activation and for treating heparanase-associated diseases or disorders such as cancer, inflammation, cardiovascular diseases, neurological diseases or viral infections. This sequence represents a heparanase inhibitor peptide used in the scope of the
                                                                                                                                                                                                                                                                                      enzyme; heparinase; vaccine; immunostimulant; cytostatic;
 New heparanase nonapeptide that
                                                                                                                                   19-MAY-2003; 2003EP-00011038
                                                                                                                                                               19-MAY-2003; 2003EP-00011038
                                                                                                                                                                                              24-NOV-2004
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                                                                                                                                                                                                                                                                                                                                Human heparanase peptide SEQ ID NO:654
                                                                                                                                                                                                                                                                                                                                                                 10-FEB-2005
                                                                                                                                                                                                                                                                                                                                                                                                                         ADU70969 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  binding protein-associated disease or disorder in a subject, a pharmaceutical composition for use in the treatment of a heparanase associated disease or disorder comprising a therapeutic amount of an agent capable of modulating heparanase activation and a pharmaceutical carrier or diluent, a method of identifying a protease activator of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Regulating heparanase activity, useful for treating heparanase-associated diseases (e.g. cancer, inflammation, cardiovascular diseases, neurological diseases or viral diseases) comprises modulating heparanase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Van-Gelder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 56; SEQ ID NO 21; 211pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             heparanase activity comprising modulating heparanase activation. The invention also relates to methods of treating a heparanase- or heparin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tissue or regulating a
                                                                                                        (DEKR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                       UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 72. 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYYLNGRTATR
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                                                         <
                                                                                       KREBSFORSCHUNGSZENTRUM.
HEIDELBERG RUPRECHT-KARLS
                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                           Beckhove
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28.4%;
72.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to a method of regulating heparanase activity in a biological process depending at least in part on
                                                         ָס
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 33.5; DB 9;
Pred. No. 1.4e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                  human leukocyte antigen; HLA;
                                                                                                                                                                                                                                                                                      immune disorder; metastasis
                                                           Sommerfeldt
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binds to a human leukocyte antigen
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RESULT 18
ADU71114
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Best Local
                        its functional derivative. A peptide of the invention has immunostimuland cytostatic activity, and is used in a vaccine. The heparinase peptis useful for preparing a medicament which induces an immune response for treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
                                                                                                                                                                                New heparanase nonapeptide that binds to a human leukocyte antigen (HLA) molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel heparanase peptide that binds to a human leukocyte antigen (HLA) molecule, where the peptide is a nonapeptide, or its functional derivative. A peptide of the invention has immunostimulant and cytostatic activity, and is used in a vaccine. The heparinase peptide is useful for preparing a medicament which induces an immune response or for treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
                                                                                                                        The invention relates
                                                                                                                                                     Disclosure; SEQ ID NO
                                                                                                                                                                                                                                                WPI; 2005-014847/02.
                                                                                                                                                                                                                                                                             Schirrmacher V,
                                                                                                                                                                                                                                                                                                            (DEKR-)
                                                                                                                                                                                                                                                                                                                                                         19-MAY-2003; 2003EP-00011038
                                                                                                                                                                                                                                                                                                                                                                                                                       24-NOV-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human heparanase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 654; 269pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
                                                                        leukocyte antigen (HLA) molecule, its functional derivative. A peptiand cytostatic activity, and is un
                                                                                                                                                                                                                                                                                                                                                                                       19-MAY-2003; 2003EP-00011038
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              heparinase;
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HEIDELBERG RUPRECHT-KARLS.
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                                                                                                                                                                                                                                                                             Beckhove P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                se; vaccine;
cytostatic;
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54.5%;
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                                                                                        to a novel heparanase peptide that binds to a hamale of the peptide is a nonapeptide.

A) molecule, where the peptide is a nonapeptide.
                                                                                                                                                   799; 269pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ
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Pred. No. 1.7e+02
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human leukocyte antigen; HLP immune disorder; metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ID NO:799
                                                                                                                                                                                                                                                                             Sommerfeldt N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A
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                                                                           The heparinase peptide
                                                                                         binds to a human
nonapeptide, or
immunostimulant
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Query Match Best Local S Matches 6

Similarity 6; Conserv

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Score 33; DB 9; Pred. No. 1.7e+02; 1; Mismatches 4

Length 15;

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                                                                                                                             Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                           95. A large number of antibody variants containing randomised peptide inserts within the complementarity determining regions (CDRs) were prepared by phage display. Libraries were subjected to eight rounds of selection to isolate variants with an antigen binding affinity at least two-fold stronger than the binding affinity of parent antibody for the target VEGF antibody. The anti-VEGF antibody variants may be useful in diagnostic assays for detecting expression of VEGF in cells, tissue or serum. They may also be used in the prevention and treatment of neoplastic diseases such as breast cancer, lung cancer and retinoblastoma, and non-neoplastic diseases including rheumatoid arthritis, psoriasis, atherosclerosis, and diabetic and other proliferarity retinoathics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibody variants with higher binding affinity than native antibodies useful for diagnosis, prevention and treatment of neoplastic and non-neoplastic diseases comprises amino acid insertion in hypervariable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; antiarthritic; antipsoriatic; antidiabetic; antiinflammatory; antiarteriosclerotic; vascular endothelial growth factor; VEGF; breast cancer; lung cancer; retinoblastoma; rheumatoid arthritis; psoriasis; atherosclerosis; diabetic retinopathy;
                                                                                                                                                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                                              proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is the randomised VH3 region of a variant of anti-
vascular endothelial growth factor (VEGF) antibody from phage library YC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 58; 110pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-387797/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Randomised VH3 sequence of anti-VEGF antibody variant Y0229-9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GETH ) GENENTECH INC.
                                                                                                                             Local Similarity
les 5; Conserv
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                                                            HYYLNN 19
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                                                                                                                                                                                                                                                              $
                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                              retinopathies
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                                                                                                                             1.
                                                                                                                                                               Score 33; DB 3;
Pred. No. 1.9e+02;
                                                                                                                                Mismatches
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                                                                                                                                                                                          Length 17;
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AAB06379 ID AAB(

AAB06379 standard; peptide; 18

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RESULT 20
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                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic; antiarthritic; antipsoriatic; antidiabetic; antiinflammatory; antiarteriosclerotic; vascular endothelial growth factor; VEGF; breast cancer; lung cancer; retinoblastoma; rheumatoid arthritis; psoriasis; atherosclerosis; diabetic retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB06381 standard; peptide;
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                                                                                                                                                                                                                                                                                                                    proliferative retinopathies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   region.
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                                                                                                                             Conservative
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71.4%;
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                                                                                                                                                        Score 33; DB 3;
Pred. No. 2.1e+02;
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                                                                                                                                                                                      Length 18;
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RESULT 22
AAB05890
ID AAB05
XX
AC AAB05
XX
DT 17-OC
XX
DE Randc
XX
KW Humar
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                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
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Humanised; antibody variant; phage display; randomised library;
cytostatic; antiarthritic; antipsoriatic; antidiabetic; antiinflammatory;
                                                 Randomised
                                                                                                                                             AAB05890 standard; peptide; 18
                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 59; 110pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antibody variants with higher binding affinity than native antibodies useful for diagnosis, prevention and treatment of neoplastic and non-neoplastic diseases comprises amino acid insertion in hypervariable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antiarteriosclerotic; vascular endothelial growth factor; VEGF; breast cancer; lung cancer; retinoblastoma; rheumatoid arthritipsoriasis; atherosclerosis; diabetic retinopathy; complementarity determining various.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Humanised; antibody variant; phage display;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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les 5; Conserv
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                                                                              (first entry)
                                              sequence of anti-VEGF antibody variant Y0261-4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence of anti-VEGF antibody variant Y0239-6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       determining region; CDR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                      28.0%;
71.4%;
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                                                                                                                                                                                                                                                                                       Score 33; DB 3;
Pred. No. 2.1e+02;
1; Mismatches 1
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                                                                                                                                                                                                                                                                                                                     Length 18;
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RESULT 23 AAB06385

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OS X KW KW KW X E X D X AC X B D I D I D

Randomised

(first entry)

AAB06385;

AAB06385 standard; peptide;

cytostatic; antiarthritic; antipsoriatic; antidiabetic; antiinflammatory; antiarteriosclerotic; vascular endothelial growth factor; VEGF; breast cancer; lung cancer; retinoblastoma; rheumatoid arthritis; psoriasis; atherosclerosis; diabetic retinopathy;

Humanised; antibody variant; phage display; randomised

VH3 sequence of anti-VEGF antibody variant Y0261-2.

Homo sapiens

complementarity

determining

region; CDR

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                                                                                                        Matches
                                                    Query Match
Best Local
                                                                                                                                                   The present sequence is the randomised VH3 region of a variant of antivascular endothelial growth factor (VEGF) antibody from phage library YC 98. A large number of antibody variants containing randomised peptide inserts within the complementarity determining regions (CDRs) were prepared by phage display. Libraries were subjected to eight rounds of selection to isolate variants with an antigen binding affinity at least two-fold stronger than the binding affinity of parent antibody for the target VEGF antibody. The anti-VEGF antibody variants may be useful in diagnostic assays for detecting expression of VEGF in cells, tissue or serum. They may also be used in the prevention and treatment of neoplastic diseases such as breast cancer, lung cancer and retinoblastoma, and non-neoplastic diseases including rheumatoid arthritis, psoriasis, atherosclerosis, and diabetic and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antibody variants with higher binding affinity than native antibodies useful for diagnosis, prevention and treatment of neoplastic and non-neoplastic diseases comprises amino acid insertion in hypervariable
                                                                                                      Sequence 18
                                                                                                                                          proliferative retinopathies
                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 60; 110pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-387797/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiarteriosclerotic; vascular endothelial growth factor; VEGF; breast cancer; lung cancer; retinoblastoma; rheumatoid arthritis; psoriasis; atherosclerosis; diabetic retinopathy; complementarity determining region; CDR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC
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                                  Similarity
5; Conserv
HYYLNNG
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                                     Conservative
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                                                     28.0%;
71.4%;
                                   Score 33; DB 3;
Pred. No. 2.1e+02;
1; Mismatches 1
                                                                    Length 18;
                                     Indels
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RESULT 24
ABB94872
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covascular endothelial growth factor (VEGF) antibody from phage library YC-
95. A large number of antibody variants containing randomised peptide
inserts within the complementarity determining regions (CDRs) were
complementarity determining regions (CDRs) were
prepared by phage display. Libraries were subjected to eight rounds of
celection to isolate variants with an antigen binding affinity at least
two-fold stronger than the binding affinity of parent antibody for the
charget VEGF antibody. The anti-VEGF antibody variants may be useful in
citarget VEGF antibody. The anti-VEGF antibody variants may be useful in
citarget VEGF antibody. The anti-VEGF antibody variants may be useful in
compositic assays for detecting expression of VEGF in cells, tissue or
serum. They may also be used in the prevention and treatment of
compositic diseases such as breast cancer, lung cancer and
cretinoblastoma, and non-neoplastic diseases including rheumatoid
carthritis, psoriasis, atherosclerosis, and diabetic and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                          Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome bladder cancer; immune response; cytotoxic T lymphocyte; human leukocyte antigen; helper T lymphocyte; HTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibody variants with higher binding affinity than native antibodies useful for diagnosis, prevention and treatment of neoplastic and non-neoplastic diseases comprises amino acid insertion in hypervariable
22-AUG-2000; 2000US-0227098P
                              22-AUG-2001; 2001WO-US026411.
                                                                                              WO200216598-A2
                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                            CTL epitope HLA peptide SEQ ID NO:401.
                                                                                                                                                                                                                                                                            17-JUN-2002
                                                                                                                                                                                                                                                                                                                                        ABB94872 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 71.55; Conservative
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Pred. No. 2.1e
1; Mismatches
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RESULT 25
ADU70797
ID ADU70
XX ADU70
AC ADU70
XX IO-FE
XX Human
XX enzym
KW inmun
XX inmun
XX EP147
XX EP147
XX EP147
XX IO-M2
PD 24-NC
XX IO-M2
PR 19-M2

10-FEB-2005

(first entry)

ADU70797;

ADU70797 standard; peptide;

9 AA

enzyme; heparinase; vaccine; human leukocyte antigen; HLA; immunostimulant; cytostatic; immune disorder; metastasis.

Human heparanase peptide SEQ ID NO:482

19-MAY-2003; 2003EP-00011038

19-MAY-2003; 2003EP-00011038.

24-NOV-2004

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YVLYSVH

8 YVLYTMH 14 Similarity 5; Conserv

Matches Query Match Best Local

Conservative

2

Mismatches

0;

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Gaps

0

27.1**%**; 71.4**%**;

Score 32; DB 5; Pred. No. 2e+06;

Length 9; Indels

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CC products in a biological sample from a patient who has or is suspected of comaring cancer. The method comprises determining the status of 158P1H4 (C) gene products in a tissue sample from an individual, comparing the status of 158P1H4 (C) comparing the status of 158P1H4 gene products in a normal sample, and comparing the status of 158P1H4 gene products in a normal sample, and comparing the presence of aberrant 158P1H4 gene products in the sample. CC identifying the presence of aberrant 158P1H4 gene products in the sample. CC production. 158P1H4 polynucleotides may be used in assessing the status of 158P1H4 gene products in normal sessed in monitoring genetic components of the malignant phenotype, in generating and characterising comparing the antibodies, for identifying agents or cellular factors that bind to 158P1H4 or its particular domain, and for generating cancer vaccines. Antibodies against 158P1H4 are useful in diagnostic and components of immunological reagents for detecting 158P1H4 cancer, in generating concer continuous to the present of language to chromosome 8q22. The concer diagnostic and concerd to chromosome 8q22. and the 158P1H4 cancer diagnostic and concerd to chromosome 8q22. ABL50400 to ABB95188 represent sequences used in the concerdiagnostic and the present invention of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monitoring 158P1H4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158P1H4 gene products in
Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       biological sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-269357/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a method for monitoring 158P1H4 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 64; Page 155; 209pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AGEN-) AGENSYS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Raitano AB,
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Page 13
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Search completed: January 26, 2006, 07:58:01 Job time: 81.1207 secs
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                                                                                                                                                        Query Match 27.1%; Score 32; DB 9; Length 9; Best Local Similarity 100.0%; Pred. No. 2e+06; Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                             The invention relates to a novel heparanase peptide that binds to a human leukocyte antigen (HLA) molecule, where the peptide is a nonapeptide, or its functional derivative. A peptide of the invention has immunostimulant and cytostatic activity, and is used in a vaccine. The heparinase peptide is useful for preparing a medicament which induces an immune response or for treating metastatic tumors. The present sequence represents a heparinase peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New heparanase nonapeptide that binds to a human leukocyte antigen (HLA) molecule or its functional derivative, useful for preparing a medicament for inducing an immune response or for treating metastatic tumors.
                                                                                                                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; SEQ ID NO 482; 269pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2005-014847/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schirrmacher V, Beckhove P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM. (UYHE-) UNIV HEIDELBERG RUPRECHT-KARLS.
                                                                                                                14 HYYLN 18
                                                                             4 HYYLN 8
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Perfect score:
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Copyright (c) 1993 - 2006 Compugen Ltd
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US-08-934-915-13
US-08-440-781-54
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US-09-440-781-7
US-09-440-781-7
US-09-287-070-9
US-09-287-070-9
US-09-480-781-85
US-09-480-781-85
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US-09-440-781-92
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13, Appl
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51, Appl
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87, Appl
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US-09-910-706A-8
PCT-US94-01238-12
US-09-99-517A-12
US-08-09-517A-12
US-08-09-517A-12
US-08-08-116-778E-7
US-08-483-528B-95
US-09-440-781-90
US-09-440-781-90
US-09-440-781-94
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US-08-934-915-13

Sequence 13, Patent No. 5

Application US/08934915

ALIGNMENTS

GENERAL INFORMATION:

APPLICANT: DILLNER, JOAKIM APPLICANT: DILLNER, LENA APPLICANT: CHENG, HWEE-MING

DILLNER, JOAKIM DILLNER, LENA

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Best Local Similarity
Watches 6; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: peptide US-08-934-915-58
                                                                                                                     US-09-440-781-44
                                                                                                                                                                                                                                     SEQ ID NO 44
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                         Sequence 44, Application US/09440781 Patent No. 6632926 GENERAL INFORMATION:
                                                  Matches
                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                     APPLICANT: Yvonne Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P14691.
CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 3.0
SOFTWARE: MICCOSOFT WORD 6.0
CURRENT APPLICATION DATA:
                                                                                                                            TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
NAME/KEY: artificial
LOCATION: 1-17
OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 19.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 813-538-3800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: LOUISE A. Foutch
REGISTRATION NUMBER: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/949,836
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                 Local Similarity nes 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: MASON & ASSOCIATES, P.A.
STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500
CITY: CLEARWATER
STATE: FLORIDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/01 FILING DATE: 22-SEP-1997
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14 HYYLNN 19
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                                                                  28.0%;
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                                                    1; Mismatches
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                                                                    Score 33; DB 2;
Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 34; DB 1;
Pred. No. 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                    0
                                                                                  Length 17;
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RESULT 2 US-08-934-915-58

Sequence 58, Application US/08934915 Patent No. 5932412

GENERAL INFORMATION:
APPLICANT: DILLNER, JOAKIM
APPLICANT: DILLNER, LENA
APPLICANT: CHENG, HWEE-MING

TITLE OF INVENTION: TITLE OF INVENTION:

INVENTION:

SYNTHETIC PEPTIDES OF HUMAN PAPILLOMAVIRUS 1, 5, 6, 8, 11, 16, 18, 31, 33 AND 56,

В Ş

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EDNVMEYVVWT-HIYL 19

Matches

Query Match

Match 37.7%; Local Similarity 62.5%;

Score 44.5; DB 1; Pred. No. 0.91;

Length 20; Indels

1;

10; Conservative 2 EXNVMXYVLYTMHYYL 17

<u>ب</u>

Mismatches

, MOLECULE TYPE: peptide US-08-934-915-13

INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
TOPOLOGY: linear

TELECOMMUNICATION INFORMATION: TELEPHONE: 813-538-3800

TELEFAX: 813-538-3820

REFERENCE/DOCKET NUMBER: NAME: LOUISE A. Foutch REGISTRATION NUMBER:

37,133

1946.6

COUNTRY: U.S.A.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 3.0
SOFTWARE: MICROSOFT WORD 6.0
CURRENT APPLICATION DATA:

CORRESPONDENCE ADDRESS:
ADDRESSIE: MASON & ASSOCIATES, P.A.
STREET: 1757 U.S. HWY. 19 NORTH, SUITE 500
CITY: CLEARWATER

FLORIDA

TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8, TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56, TITLE OF INVENTION: USEFUL IN IMMUNOASSAY FOR TITLE OF INVENTION: DIAGNOSTIC PURPOSES NUMBER OF SEQUENCES: 193

APPLICATION NUMBER: US/08/934,915
FILING DATE: 22-SEP-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/949,836

ATTORNEY/AGENT INFORMATION:

FILING DATE:

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RESULT 6
US-09-440-781-57
; Sequence 57, Applicati
; Patent No. 6633926
; GENERAL INFORMATION:
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; NAME/KEY: artificial
; LOCATION: 1-18
; OTHER INFORMATION: variant CDR sequence
US-09-440-781-51
                                                                                                                                                                                                                                                                                                                                                    US-09-440-781-53
                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Yvonne Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
SEQ ID NOS: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 28.0
Best Local Similarity 71.4
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 51
LENGTH: 18
TYPE: PRT
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GENERAL INFORMATION:
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Patent No. 6632926
APPLICANT: Yvonne Man-yee Chen et al. TITLE OF INVENTION: ANTIBODY VARIANTS FILE REFERENCE: P1469R1 CURRENT APPLICATION NUMBER: US/09/440,781 CURRENT FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Yvonne Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: artificial LOCATION: 1-18
OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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                                                                                                                        Application US/09440781
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                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                              28.0%;
71.4%;
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Pred. No. 62;
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; SEQ ID NO 65
; LENGTH: 17
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; NAME/KEY: artificial
; LOCATION: 1-17
; OTHER INFORMATION: variant CDR sequence
US-09-440-781-65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 87

LENGTH: 18

TYPE: PRT

ORGANISM: artificial sequence
FEATURE:
NAME/KEY: artificial
LOCATION: 1-18
OTHER INFORMATION: variant CDR sequence
US-09-440-781-87
                                                                                                                                                                                                                                                                                                                                 RESULT 8
US-09-440-781-65
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Sequence 87, Application US/09440781
Patent No. 6632926
GENERAL INFORMATION:
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NAME/KEY: artificial
LOCATION: 1-18
OTHER INFORMATION: variant CDR sequence
US-09-440-781-57
                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Yvonne Man-yee Chen et al.
                                                                                                                                                                                                                                                                                                  Sequence 65, Application US/09440781 Patent No. 6632926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 99
SEQ ID NO 57
LENGTH: 18
TYPE: PRT
             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
                                                                                                                                                                                                                                  TITLE OF INVENTION: ANTIBODY VARIANTS FILE REFERENCE: P1469R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Yvonne Man-yee Chen et al. TITLE OF INVENTION: ANTIBODY VARIANTS FILE REFERENCE: P1469R1
CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 28.0%;
Local Similarity 71.4%;
es 5; Conservation
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3 HYYLKDG 9
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5; Conserv
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71.4%;
Score 32;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 33; DB
Pred. No. 62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 33; DB
Pred. No. 62;
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 85;
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                2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                Length 17;
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Local Similarity

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US-10-395-434A-2
Sequence 2, Application US/10395434A
Patent No. 6939948
GENERAL INFORMATION:
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                                                               RESULT 11
                                                                                                                                                                                                                  ; OTHER INFORMATION: variant CDR sequence US-09-440-781-35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 10
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                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: YVONDE MAD-YEE Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 99
SEQ ID NO 35
LENGTH: 18
TYPE: PRT
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                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 35, Application US/09440781 Patent No. 6632926
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                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
APPLICANT: Ferrone, Soldano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Yvonne Man-yee Chen et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 17
TYPE: PRT
ORGANIAN: artificial sequence
FEATURE:
NAME/KEY: artificial
LOCATION: 1-17
OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: artificial
LOCATION: 1-18
                                                                                                                                                                                                                                                                                   ORGANISM: artificial
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                                                                                                                                 14 HYYLNNG 20
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                                                                                                                                                            Similarity 5; Conserv
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                                                                                                                                                                         27.1%;
71.4%;
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Pred. No. 91;
0; Mismatches
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85;
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Patent No. 6632926
GENERAL INFORMATION:
APPLICANT: YVONDE MAN-Yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/395,434A
CURRENT FILING DATE: 2003-03-24
PRIOR APPLICATION NUMBER: US 60/366,558
PRIOR FILING DATE: 2002-03-22
NUMBER OF SEQ ID NOS: 9
SEQ ID NO 2
                                                                                                                                                                                                                                                               Sequence 7, Application US/08934915 Patent No. 5932412
                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: GD2 Peptide Mimics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 03551.009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: artificial
LOCATION: 1-17
OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: J8C peptide mimic for GD2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                  APPLICANT: DILLNER, JOAKIM
APPLICANT: DILLNER, LENA
APPLICANT: CHENG, HWEE-MING
TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,
TITLE OF INVENTION: 11, 16, 31, 33 AND 56,
TITLE OF INVENTION: USEFUL IN IMMUNOASSAY FOR
TITLE OF INVENTION: DIAGNOSTIC PURPOSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 12
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                                                                 CORRESPONDENCE ADDRESS:
                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity les 4; Conserv
                              ADDRESSEE: MASON & ASSOCIATES, P.J. STREET: 17757 U.S. HWY. 19 NORTH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      14 HYYLNN 19
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                  CLEARWATER
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FLORIDA
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                                                                                   11, 16, 18, 31, 33 AND 56, USEFUL IN IMMUNOASSAY FOR DIAGNOSTIC PURPOSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                         25.4%;
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Pred. No. 1.8e+02;
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Pred. No. 83;
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                                  SUITE 500
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COMPUTER FRADABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC OPERATING SYSTEM:

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RESULT 15
US-08-370-909-10
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                                                                                                                                                                                                                            ; LENGTH: 20
; TYPE: PRT
; ORGANISM: Pneumococcus
US-09-287-070-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 25.4%;
Best Local Similarity 50.0%;
Matches 6; Conservative
                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                       SEQ ID NO 9
                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                   APPLICANT: Tuomanen, Elaine
APPLICANT: Gosink, Khoosheh
APPLICANT: Masure, Robert
TITLE OF INVENTION: IDENTIFICATION AND CHARACTERIZATION OF NOVEL
TITLE OF INVENTION: DIACNOSTIC AND THERAPEUTIC USES THEREOF
FILE OF INVENTION: DIACNOSTIC AND THERAPEUTIC USES THEREOF
FILE REFERENCE: 1340-1-024cip
CURRENT APPLICATION NUMBER: US/09/287,070A
CURRENT FILING DATE: 1999-04-06
EARLIER FILING DATE: 1999-04-06
EARLIER APPLICATION NUMBER: 09/196,389
EARLIER FILING DATE: 1998-11-19
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 20 amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: LOUISE A. Foutch
REGISTRATION NUMBER: 37,133
REFERENCE/DOCKET NUMBER: 1946.6
TELECOMMUNICATION INFORMATION:
TELEPHONE: 813-538-3800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/949,836
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: Windows 3.0 SOFTWARE: Microsoft Word 6.0 CURRENT APPLICATION DATA:
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FILING DATE: 22-SEP-1997
CLASSIFICATION: 435
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                                                                                                                                                    7;
                                                                                                                                                                      Similarity
                                                                         YYLNANGA 17
                                                                                                                                                    Conservative
                                                                                                                                                                    25.0%;
87.5%;
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                                                                                                                                                                      Score 29.5; DB 2;
Pred. No. 2.7e+02;
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Pred. No. 2.
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                                                                                                                                                                    2.7e+02;
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2.2e+02;
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; LENGTH: 9
; TYPE: AMINO ACID
; STRANDEDNESS: UNKNOWN
; TOPOLOGY: UNKNOWN
; MOLECULE TYPE: PEPTIDE
US-08-370-909-10
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US-09-774-639-201
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                                                                                                                                APPLICANT: Rosen et al.
TITLE OF INVENTION: 90 Human Secreted Proteins
FILE REFERENCE: PZ013P1
CURRENT APPLICATION NUMBER: US/09/774,639
CURRENT FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/244,112
PRIOR FILING DATE: EARLIER FILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 371
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 201
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                   Sequence 201, Application Patent No. 6806351
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 24.6%;
Best Local Similarity 66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (212) 751-684
TELEX: 421792
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: ROBBINS, PAUL F.; ROSENBERG,
APPLICANT: STEVEN A.
APPLICANT: STEVEN A.
TITLE OF INVENTION: P15 AND TYROSINASE
TITLE OF INVENTION: AND THERAPEUTIC MET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: CAROL M. GRUPPI
REGISTRATION NUMBER: 37,341
REFERENCE/DOCKET NUMBER: 2026
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
NAME/KEY: SITE
LOCATION: (9)
OTHER INFORMATION: Xaa equals stop translation
                                                                  ORGANISM: Homo sapiens FEATURE:
                                                                                                        TYPE: PRT
                                                                                                                         ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/370,909
FILING DATE: 10-JAN-1995
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
STREET: 345 PARK AVENUE
CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 10154
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4; Conservative
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Pred. No. 4.6e+05;
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Query Match
Best Local Similarity
"hes 5; Conserva
                                                                               ; LENGTH: 15
; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Description
US-08-793-331-13
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US-09-419-381-14
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US-08-793-331-13
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SEQ ID NO 14
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6071877
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Best Local 9
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Best Local :
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EARLIER FILING DATE: 1995-08-24
EARLIER APPLICATION NUMBER: FR 94/10299
EARLIER FILING DATE: 1994-08-25
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                   APPLICANT: DELECLUSE, ARMELLE
APPLICANT: THIERY, ISABELLE
TITLE OF INVENTION: NEW POLYPEPTIDES HAVING A TOXIC ACTIVITY AGAINST
TITLE OF INVENTION: INSECTS OF THE DIFTERAE FAMILY
FILE REFERENCE: 0660-0116-0 PCT
CURRENT PEPLICATION NUMBER: US/08/793,331
CURRENT FILING DATE: 1997-05-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/99/419,381
CURRENT FILING DATE: 1999-10-15
PRIOR APPLICATION NUMBER: 60/104,465
PRIOR FILING DATE: 1998-10-16
NUMBER OF SEQ ID NOS: 122
COPTUBARY: 05-50-51
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APPLICANT: Rozinov, Michael N.
TITLE OF INVENTION: Fluorescent Dye Binding Peptides
FILE REFERENCE: A65681-1/DJB/RMS/DSS
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                 Conservative
                                                                                                  Description of Artificial Sequence: JEG66
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               Score 29; DB
Pred. No. 2.3
0; Mismatches
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Pred. No. 1.8e+02
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Pred. No. 4.6e+05;
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APPLICANT: Yvonne Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
LENGTH: 17
TYPE: PRT
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Best Local Similarity
Thes 4; Conserve
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US-09-440-781-49
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                                                                                              RESULT 21
US-09-440-781-92
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Sequence 92, Application US/09440781 Patent No. 6632926
GENERAL INFORMATION:
APPLICANT: Yvonne Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
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LENGTH: 17
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Yvonne Man-yee Chen et al. TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PEATURE: NAME/KEY: artificial
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                                                                                                                                                                                                                                                                                                                  NAME/KEY: artificial LOCATION: 1-17
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80.0%;
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Pred. No. 2.7e+02;
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Pred. No. 2.7e+02;
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US-09-461-325-279
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                                                                                                                                                            RESULT 23
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NAME/KEY: artificial
; LOCATION: 1-17
; OTHER INFORMATION: variant CDR sequence
US-09-440-781-92
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Sequence 279, Application US/10012542
Patent NO. 6627741
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 94 Human Secreted ProffILE REFERENCE: PZ02991
CURRENT APPLICATION UNMERR: US/10/012,542
CURRENT FILING DATE: 2001-12-12
                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 5; Conserv
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NUMBER OF SEQ ID NOS: 532
SOFTWARE: PALENTIN VEY. 2.0
SEQ ID NO 279
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Best Local
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GENERAL INFORMATION:
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CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
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EARLIER APPLICATION NUMBER: 60/090,113
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EARLIER FILING DATE: 1998-06-16
EARLIER APPLICATION NUMBER: 60/089,508
EARLIER FILING DATE: 1998-06-16
EARLIER APPLICATION NUMBER: 60/089,509
EARLIER FILING DATE: 1998-06-16
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CURRENT APPLICATION NUMBER: US/09/461,325A
CURRENT FILING DATE: 1999-12-14
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TITLE OF INVENTION: 94 Human Secreted Proteins
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TYPE: PRT
ORGANISM: artificial sequence
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                                                      94 Human Secreted Proteins
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Pred. No. 2.7e+02
1; Mismatches
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Pred. No. 2
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-115-123-279
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-012-542-279
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                                       Query Match
Best Local Similarity barrance
Thes 5; Conservative
                                                                                                                                                                                                        SEQ ID NO 279
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence 279, App
Patent No. 677421
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Best Local Similarity
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TITLE OF INVENTION: 94 Human Secreted Pro
FILE REFERNCE: PSC029G30APID2
CURRENT APPLICATION NUMBER: US/10/115,123
CURRENT FILING DATE: 2002-04-04
                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/090,113
PRIOR FILING DATE: 1998-06-22
NUMBER OF SEQ ID NOS: 532
SOFTWARE: Patentin Ver. 2.0
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PRIOR FILING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: 60/089,508
PRIOR FILING DATE: 1998-06-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT/US99/13418 PRIOR FILING DATE: 1999-06-15
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                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/089,510 FILING DATE: 1998-06-16
                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/089,509 FILING DATE: 1998-06-16
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                                12 TMHYYLNN 19
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TVLFYLNN
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                                                                                 Score 29; DB 2;
Pred. No. 2.8e+02;
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Pred. No. 2.8e+02
                                                                  Mismatches
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APPLICATION NUMBER:
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APPLICATION NUMBER:
DATE: 1998-06-16
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APPLICATION NUMBER: 60/090,113
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                                                                                                   Length 18;
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US-09-178-093B-19

Sequence 19, Application US/09178093B

Patent No. 6660846

GENERAL INFORMATION:
APPLICANT: Richard J. Reimer
APPLICANT: Richard J. Reimer
APPLICANT: Erik M. Jorgenson
APPLICANT: Erik M. Jorgenson
APPLICANT: Erik M. Jorgenson
APPLICANT: Kim Schuske
ITTLE OF INVENTION: Composition and Method
FILE REFERENCE: 2002-005.30
CURRENT FILING DATE: 2001-08-20
PRIOR APPLICATION NUMBER: US/09/178,093B
CURRENT FILING DATE: 2001-08-20
PRIOR APPLICATION NUMBER: 60/03,012
PRIOR APPLICATION N
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Perfect score:
Sequence:
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Maximum
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Match
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Maximum
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128.563 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US11A_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Copyright (c) 1993 - 2006 Compugen
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 US-10-346-198-43
US-09-880-748-2216
US-10-293-418-2216
US-10-467-758-35
US-09-880-748-2259
US-10-086-814-30
US-10-086-814-30
US-10-086-814-36
US-10-624-153-51
US-10-624-153-51
US-10-624-153-87
US-10-624-153-87
US-10-624-153-87
US-10-624-153-87
US-10-293-418-2909
US-11-094-60-104
US-10-624-153-86
US-09-880-748-2968
US-09-880-748-2968
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 2216, Appl
2216, Appl
35, Appl
35, Appl
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10, Appl
10, Appl
11, Appl
21, Appl
51, Appl
     30
       US-10-293-418-2158
US-10-293-418-299
US-10-293-418-299
US-10-293-418-291
US-09-880-748-2217
US-09-880-748-2217
US-09-880-748-2217
US-09-880-748-2219
US-09-880-748-2219
US-10-293-418-2219
US-10-245-871-489
US-10-245-871-489
US-10-245-871-512
US-10-245-871-512
US-10-245-871-512
US-10-245-871-512
US-10-245-871-513
US-10-253-286-112
US-10-341-201-29
US-09-748-875-33
US-10-341-201-29
US-09-298-5238-33
US-10-341-201-31
US-10-341-201-31
US-10-341-201-31
US-10-341-201-31
US-10-341-201-33
US-10-341-201-31
US-10-341-201-33
US
IS-09-774-639-201

IS-10-105-232-169

IS-10-189-437-156

IS-10-860-050-169

IS-10-692-151-14

IS-10-692-151-14

IS-10-690-18

IS-09-880-748-2258

IS-09-880-748-2258

IS-09-880-748-2258
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     2158 Ap 2293, Ap 2291, Ap 2291
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ALIGNMENTS

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PRIOR APPLICATION NUMBER: 2003-01-16;
PRIOR APPLICATION NUMBER: 60/337,409
PRIOR FILING DATE: 2002-05-01
NUMBER OF SEQ ID NOS: 1149
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 43
LENGTH: 20
TYPE: PRT
ORGANISM: Oryza sativa
US-10-346-198-43
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US-10-346-198-43
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                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-03-25
PRIOR FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 3239
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2216, Application US/09880748 Publication No. US20030059937A1
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Publication No. US20040043485A1
                                                                                           Query Match 29.7%;
Best Local Similarity 71.4%;
Matches 5; Conservative
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Best Local Similarity 63.6%;
                                                                                                                                                                            LENGTH: 16
TYPE: PRT
ORGANISM: Homo sapiens
-09-880-748-2216
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APPLICANT: JIANG, NING
APPLICANT: BAO, ZHRONG
APPLICANT: ZHANG, XIAOYU
APPLICANT: EDDY, SEAN R.
TITLE OF INVENTION: TRANSPOSABLE ELEMENTS IN RICE AND METHODS OF USE
FILE REFERENCE: 18465-0018
CURRENT APPLICATION NUMBER: US/10/346,198
CURRENT FILING DATE: 2003-01-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
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                 YVLHYYL 15
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Pred. No. 1.3e+02;
                                                                                                                 Score 35; DB 3;
Pred. No. 1.5e+02;
                                                                                                  Mismatches
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APPLICANT: Hupp, Theodore
APPLICANT: Hupp, Theodore
APPLICANT: Dornan, David
ITITLE OF INVENTION: Screening Method and Agents
FILE REFERENCE: 9013.54
CURRENT APPLICATION NUMBER: US/10/467,758
CURRENT FILING DATE: 2003-08-13
PRIOR APPLICATION NUMBER: DCT/GB02/00640
PRIOR PILING DATE: 2002-02-13
PRIOR APPLICATION NUMBER: GB 0103508.8
PRIOR FILING DATE: 2001-02-13
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; ORGANISM: Homo sapiens
US-10-293-418-2216
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CURRENT FILING DATE: 2002-11-27
PRIOR APPLICATION UNMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR PELING DATE: 2001-12-19
PRIOR PELING DATE: 2001-06-15
PRIOR PELING DATE: 2001-06-15
PRIOR PELING DATE: 2001-05-25
PRIOR PILING DATE: 2001-05-25
PRIOR PILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR PILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/277,248
PRIOR PILING DATE: 2001-03-16
PRIOR PILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                   US-10-467-758-35
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PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2216
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                                                                                                                                                                                                                                  SEQ ID NO 35
LENGTH: 12
                                                                    Query Match 28.8%;
Best Local Similarity 50.0%;
Matches 6; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-02-13
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin version 3.1
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                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
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11 YTMHYYLNNGAT 22
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                                                                                               Score 34; DB 4; 1
Pred. No. 1.5e+02;
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                                                                         Mismatches
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US-09-880-748-2259
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Best Local Similarity 83...
Conservative
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PRIOR FILING DATE: 2000-06-15
PRIOR PELICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR PELICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
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SEQ ID NO 2259
LENGTH: 16
                                                                                   PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2259
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GENERAL INFORMATION
ORGANISM: Homo sapiens
10-293-418-2259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS FILE REFERENCE: PF523P2 CURRENT APPLICATION NUMBER: US/10/293,418 CURRENT FILING DATE: 2002-11-27
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                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/340,817 PRIOR FILING DATE: 2001-12-19
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PRIOR FILING DATE: 2001-11-16
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                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/277,379
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o. US20030223996A1
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RESULT 8
US-10-086-814-36
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 36
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Publication No.
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SEQ ID NO 30
LENGTH: 21
TYPE: PRT
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Best Local Similarity
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APPLICANT: Olson, William C.
TITLE OF INVENTION: SULFATED CCR5 PEPTIDES FOR HIV-1 INFECTION
                                                                                                                                                                                                                               FILE REFERENCE: 61010-AB-1
CURRENT APPLICATION NUMBER: US/10/086,814
CURRENT FILING DATE: 2002-02-28
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APPLICANT: Olson, Wlliam C.
                                                                                                               LENGTH: 21
TYPE: PRT
ORGANISM: Homo sapiens
                                               NAME/KEY: MOD_RES
LOCATION: (9)...(9)
OTHER INFORMATION: PHOSPHORYLATION
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LOCATION: (21)..(21)
OTHER INFORMATION: BIOTIN
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LOCATION: (13)..(13)
OTHER INFORMATION: SULFATATION
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NAME/KEY: MOD RES
LOCATION: (9)...(9)
OTHER INFORMATION: SULFATATION
FEATURE:
NAME/KEY: MOD_RES
LOCATION: (13)..(13)
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Local Similarity 41.7%;
hes 5; Conservativo
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83.3%;
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Pred. No. 2.8e+02;
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OTHER INFORMATION: PHOSPHORYLATION

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RESULT 10
US-10-722-502-10
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SEQ ID NO 147622
LENGTH: 21
TYPE: PRT
ORGANISM: Glycine max
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         SOFTWARE: PatentIn version 3.2
SEQ ID NO 10
LENGTH: 14
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Publication No. US20040170631A1
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Best Local Similarity
Matches 5; Conserv
                                                          APPLICANT: Feinstein, Elena
APPLICANT: Van Gelder, Joel M.
APPLICANT: Vlodavsky, Israel
APPLICANT: Priedmann, Yael
TITLE OF INVENTION: HEPARANASE ACTIVITY NEUTRALIZING ANTI- HEPARANASE MONOCLONAL
TITLE OF INVENTION: ANTIBODY AND OTHER ANTI-HEPARANASE ANTIBODIES
FILE REFERENCE: 26872
CURRENT APPLICATION UNMER: US/10/722,502
CURRENT FILING DATE: 2003-11-28
MUMBER OF SEQ ID NOS: 11
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APPLICANT: Peretz, Tuvia
APPLICANT: Miron, Daphna
APPLICANT: Shlomi, Yinon
APPLICANT: Pecker, Iris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53223) B
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CURRENT FILING DATE: 2003-04-28
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APPLICANT: Kovalic David K
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TYPE: PRT
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Pred. No. 2.8e+02
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Pred. No. 2.8e+02
4; Mismatches
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US-10-916-598-21
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                                                                               Sequence 21, Application US/10916598
Publication No. U320050042213A1
GENERAL INFORMATION:
APPLICANT: Van Gelder, Joel M.
APPLICANT: Miron, Daphna
TITLE OF INVENTION: 27413
FILE REFERENCE: METHODS AND PHARMACEUTICAL COMPOSITIONS FOR MODULATING HEPARANASE
ACTIVATION AND USES THEREOF
CURRENT APPLICATION NUMBER: US/10/916,598
CURRENT APPLICATION UNMBER: US/10/916,598
CURRENT FILING DATE: 2004-08-12
NUMBER OF SEQ ID NOS: 37
SOFTWARE: PACENTIN version 3.2
SEQ ID NO 21
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 26128
CURRENT APPLICATION NUMBER: US/10/645,659
CURRENT FILING DATE: 2003-08-22
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin version 3.2
SEQ ID NO 10
LENGTH: 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 8; Conserv
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: HEPARANASE ACTIVITY NEUTRALIZING ANTI- HEPARANASE MONOCLONAL TITLE OF INVENTION: ANTIBODY AND OTHER ANTI-HEPARANASE ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                               LENGTH: 15
TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial sequence
            OTHER INFORMATION: Synthetic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE: OTHER INFORMATION: Functional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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les 8; Conservative
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Miron, Daphna
Shlomi, Yinon
Pecker, Iris
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Friedmann, Yael
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72.7%;
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Pred. No. 2.2e
0; Mismatches
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Pred. No. 2.2e+02;
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Query Match
Best Local Similarity
Thes 5; Conserv
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; Sequence 44, Application US/10624153
; Publication No. US20040086502A1
                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 51
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 51, Application US/10624153 Publication No. US20040086502A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR FILING DATE: 1999-11-16
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CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
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APPLICANT: LOWMAN, HENRY
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                                                                                                               FILE REFERENCE: P1469R1C:
                                                                                                                                                                                                                                                                                                                             APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: CHEN, APPLICANT: LOWMA
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EQ ID NO 44
   NAME/KEY: artificial LOCATION: 1-18
                                       FEATURE:
OTHER INFORMATION: sequence is synthesized
                                                                                           ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: 1-17
OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: artificial
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OTHER INFORMATION: sequence is synthesized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: artificial sequence
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83.3%;
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72.7%;
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Pred. No. 3.2e+02;
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Pred. No. 2.
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2.3e+02;
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                                                                                     APPLICANT: CHEN, YVONNE M.
APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1C1
CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1999-11-16
PRIOR FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 57
                                                                                                                                                                                                                                                                                                                                                             US-10-624-153-57
; Sequence 57, Application US/10624153
; Publication No. US20040086502A1
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Best Local Similarity 71.4
Matches, 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
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APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: CHEN, YVONNE M. APPLICANT: LOWMAN, HENRY
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OTHER INFORMATION: variant CDR sequence
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OTHER INFORMATION: sequence is synthesized
FEATURE:
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OTHER INFORMATION: sequence is synthesized
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TYPE: PRT
                                     ORGANISM: artificial sequence
                                                                           ENGTH: 18
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71.4%;
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Pred. No. 3.4e+02;
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Pred. No. 3.4e+02;
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CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 87
LENGTH: 18
TYPE: PRI
TYPE: PRI
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                                                                                                                                                                                                                                                                                          Sequence 401, Application US/0935384 Publication No. US20030166526A1 GENERAL INFORMATION:
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Best Local Similarity 71.4
Matches 5; Conservative
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1C1
            APPLICANT: LEVIN, ELANA
APPLICANT: FARIS, MARY
APPLICANT: GE, WANGWAO
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158-20033.00
CURRENT FILING DATE: 2001-08-22
CURRENT FILING DATE: 2001-08-22
CURRENT FILING DATE: 2001-08-22
                                                                                                                                                                                           APPLICANT: CHALLITA-EID, PIA
APPLICANT: HUBERT, RENE
APPLICANT: RAITANO, ARTHUR
APPLICANT: AFAR, DANIEL
APPLICANT: LEVIN, ELANA
PRIOR APPLICATION NUMBER: 60/227,098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PEATURE:
NAME/KEY: artificial
LOCATION: 1-18
OTHER INFORMATION: variant CDR sequence
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NAME/KEY: artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: sequence is synthesized
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71.4%;
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OF BLADDER AND
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Sequence 2909, Application US/10293418

Publication No. US20030223996A1

GENERAL INFORMATION:
APPLICANT: Ruben et al.

ITITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P2

CURRENT APPLICATION NUMBER: US/10/293,418

CURRENT FILING DATE: 2002-11-27

PRIOR APPLICATION NUMBER: 60/331,469

PRIOR FILING DATE: 2001-11-16

PRIOR APPLICATION NUMBER: 60/340,817

PRIOR APPLICATION NUMBER: 60/340,817
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CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/21,210
PRIOR FILING DATE: 2000-06-5
PRIOR FILING DATE: 2000-10-17
PRIOR PPLICATION NUMBER: 60/276,248
PRIOR PPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-21
PRIOR PILING DATE: 2001-03-21
PRIOR PILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 3239
SOFTWARE: Patentin Ver. 2.0
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US-10-293-418-2909
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Matches
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523
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TYPE: PRT
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FILING DATE:
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2001-05-25
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Pred. No. 4e+02;
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Pred. No. 1.7e+06;
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FRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-6-16
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2909
LENGTH: 15
TYPE: PRT
ORGANISM: Homo Bapiens
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; Sequence 65, Application US/10624153
; Publication No. US20040086502A1
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                                                           GENERAL INFORMATION:
APPLICANT: CHEN, YVONNE M.
APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 104
LENGTH: 15
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Matches 6; Conserv
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Matches 4; Conservative
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CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: CARTER, Graham
TITLE OF INVENTION: METHOD FOR MAPPING AND ELIMINATING
TITLE OF INVENTION: T-CELL EPITOPES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Potential T-cell Epitopes
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75.0%;
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Pred. No. 4e+02;
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RESULT 24
US-10-624-153-35
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US-10-624-153-84
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APPLICANT: MULLER, YVES

TITLE OF INVENTION: ANTIBODY VARIANTS

FILE REFERENCE: P1469R1C1

CURRENT APPLICATION NUMBER: US/10/624,153

CURRENT FILING DATE: 2003-07-21

PRIOR APPLICATION NUMBER: US 09/440,781

PRIOR FILING DATE: 1999-11-16

PRIOR FILING DATE: 1999-11-18

PRIOR FILING DATE: 1998-11-18

NUMBER OF SEQ ID NOS: 99
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LENGTH: 17
TYPE: PRT
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 84
           Sequence 35, App
Publication No.
                                                                                                                                                                             Matches
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Best Local Similarity
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PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
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APPLICANT: LOWM
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REATURE:
NAME/KEY: artificial
LOCATION: 1-17
OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                                                                                                                                                                        LENGTH: 17
TYPE: PRT
ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                            FEATURE:
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             Application US/10624153
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71.4%;
                                                                                                                                                                                           27.1%; Score 32; 100.0%; Pred. No.
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Pred. No. 4.6e+02;
0; Mismatches
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                                                                                                                                                                                                          DB 4;
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                                                                                                                                                                                              4.6e+02
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APPLICANT: CHEN, YVONNE M.
APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS

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P1469R1C1

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US-09-880-748-2158
Sequence 2158, Application US/09880748
| Publication No. US20030059937A1
| GENERAL INFORMATION:
| APPLICANT: Ruben et al.
| TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
| FILE REFERENCE: pF523
| CURRENT APPLICATION NUMBER: US/09/880,748
| CURRENT FILING DATE: 2001-06-15
| PRIOR APPLICATION NUMBER: 60/212,210
| PRIOR APPLICATION NUMBER: 60/240,816
| PRIOR APPLICATION NUMBER: 60/276,248
| PRIOR APPLICATION NUMBER: 60/276,248
| PRIOR APPLICATION NUMBER: 60/277,379
| PRIOR APPLICATION NUMBER: 60/293,499
| PRIOR APPLICATION 
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CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 35
LENGTH: 18
TYPE: PRT
ORGANIEM: artificial sequence
FEATURE:
OTHER INFORMATION: sequence is synthesized
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NAME/KEY: artificial
LOCATION: 1-18
TOTHER INFORMATION: variant CDR sequence
US-10-624-153-35
Search completed: January 26, 2006, 08:38:36
Job time : 75.75 secs
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Best Local Similarity 71.4%;
Matches 5; Conservative
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Best Local Similarity 71.4%;
Matches 5; Conservative
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2: /cgn2_6/ptodata/1/pubpaa/USOS_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*

8: /cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*
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US-10-939-890-205
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2216 Ap
255, App
256, App
256, App
257, App
75, App
75, App
75, App
76, App
2003 Ap
2014, Ap
2211, Ap
2211, Ap
2211, App
211, App
213, App
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    US-11-054-515-226
US-11-054-515-226
US-11-054-515-230
US-11-054-515-237
US-11-054-515-237
US-11-054-515-233
US-11-054-515-233
US-11-054-515-233
US-11-054-515-233
US-11-010-748A-73
US-11-010-748A-84
US-11-010-748A-13
US-11-010-748A-13
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US-11-054-515-212
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US-11-054-515-223
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RESULT 2
US-11-054-515-2259
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PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
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Best Local S
Matches 5
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SEQ ID NO 2216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2216, Application US/11054515
Publication No. US20050255532A1
GENERAL INFORMATION:
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TITLE OF INVENTION: Antibodies that Immunospecifically
FILE REFERENCE: PF523P3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
                                                                                             PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-1
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
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PRIOR
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CURRENT APPLICATION NUMBER: US/11/054,515
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TITLE OF INVENTION: Ant
                                                       PRIOR APPLICATION NUMBER: 10/293,418 PRIOR FILING DATE: 2002-11-14
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                  APPLICATION NUMBER: 60/331,469 FILING DATE: 2001-11-16
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APPLICATION NUMBER: 60/340,817
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; Pred. No. 5.3;
1; Mismatches
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US-11-045-024-10532
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Sequence 10532, A
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RESULT 4
US-10-511-559-256
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CURRENT APPLICATION NUMBER: US/10/511,559
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR PFLING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR FILING DATE: 2002-04-18
PRIOR FILING DATE: 2002-04-18
PRIOR APPLICATION NUMBER: EP 03006554.4
PRIOR FILING DATE: 2003-03-24
NUMBER OF SEQ ID NOS: 1147
SOFTWARE: FastSEQ for Windows Version 4.0
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; LENGTH: 16
; TYPE: PRT
                                                                                                                                                                                                                                                                                    SEQ ID NO 255
LENGTH: 13
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Best Local :
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Best Local Similarity
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APPLICANT: BAKER, Matthew
APPLICANT: CARR, Francis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
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Remaining Prior Application of NUMBER OF SEQ ID NOS: 3247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: MODIFIED FACTOR VIII
                                                                                                                                                                                                              FEATURE: OTHER INFORMATION: Potential Epitope of human Factor VIII
                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                        TYPE: PRT
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les 5; Conserv
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APPLICATION NUMBER: 09/880,748
FILING DATE: 2001-06-15
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o. US20050256304A1
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Pred. No. 7.6;
0; Mismatches
                                                                                                                                                   Score 32; DE
Pred. No. 13;
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GENERAL INFORMATION:
APPLICANT: JONES, Tim
APPLICANT: BAKER, Mat
APPLICANT: CARR, Fran

BAKER, Matthew CARR, Francis,

Sequence 256, Appropriate Publication No.

Application US/10511559 o. US20050256304A1

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RESULT 6
US-10-511-559-75
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US-10-511-559-257
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CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR FILING DATE: 2002-04-18
PRIOR FILING DATE: 2002-04-18
PRIOR FILING DATE: 2003-03-24
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US-10-511-559-256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/511,559
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR FILING DATE: 2002-04-18
PRIOR APPLICATION NUMBER: EP 03006554.4
PRIOR FILING DATE: 2003-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 1147
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 257
LENGTH: 13
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 256
                                       Sequence 75, Application US/10511559
Publication No. US20050256304A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                      Query Match 27.1%;
Best Local Similarity 75.0%;
Matches 6; Conservative
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Best Local :
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APPLICANT: JONES, Tim APPLICANT: BAKER, Matthew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: JONES, Tim
APPLICANT: BAKER, Mat
APPLICANT: CARR, Fran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: MODIFIED FACTOR VIII FILE REFERENCE: MER-133
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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5. US20050256304A1
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Pred. No. 13;
0; Mismatches 2; Indels
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PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR FILING DATE: 2002-04-18
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-24
NUMBER OF SEQ ID NOS: 1147
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 75
LENGTH: 15
                                                                                        ; ORGANISM: Homo sapiens
US-11-054-515-2909
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                                                                                                                                                                                                                                         PRIOR FILLING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR APPLICATION NUMBER: 09/880,748
PRIOR APPLICATION NUMBER: 09/880,748
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-10-17
                                                                                                                                Remaining Prior Application
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2909
LENGTH: 15
TYPE: PRT
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  Matches
                     Query Match
Best Local Similarity
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Best Local S
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
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TITLE OF INVENTION: MODIFIED FACTOR VIII
FILE REFERENCE: MER-133
CURRENT APPLICATION NUMBER: US/10/511,559
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR FILING DATE: 2003-04-17
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PRIOR TILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/331,469
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Similarity 75.0%;
6; Conservative
Conservative
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                     27.1%;
33.3%;
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Pred. No. 15;
0; Mismatches
4; Mismatches
                     Score 32; DB
Pred. No. 15;
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PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-19
PRIOR FILING DATE: 2001-12-19
                                                                                                                                                                                                                                                                                      Sequence 2896, Application US/11054515
Publication No. US2005025532A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3
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NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2158
LENGTH: 20
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GENERAL INFORMATION:
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Best Local
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CURRENT FILLING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
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CURRENT FILING DATE: 2005-02-10
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TITLE OF INVENTION: Ant
FILE REFERENCE: PF523P3
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                                         APPLICATION NUMBER: 10/293,418
FILING DATE: 2002-11-14
APPLICATION NUMBER: 60/31,469
FILING DATE: 2001-11-16
APPLICATION NUMBER: 60/340,817
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FILING DATE: 2001-05-25
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FILING DATE: 2001-12-19
APPLICATION NUMBER: 09/880,748
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71.4%;
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LENGTH: 20
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LENGTH: 20
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PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
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PRIOR TILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
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TITLE OF INVENTION: Ant
FILE REFERENCE: PF523P3
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PRIOR FILING DATE: 2000-10-17
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PRIOR TILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
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PRIOR FILING DATE: 2001-12-19
                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
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PRIOR FILING DATE: 2001-06-15
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nes 5; Conserv
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                                   15 YYLNNGA 21
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YYIGNGA 17
                                                                      Conservative
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                                                                                     27.1%;
71.4%;
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71.4%;
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                                                                      Score 32; DB
Pred. No. 20;
1; Mismatches
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Pred. No.
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CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR APPLICATION NUMBER: 60/331,469
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CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT FILING DATE: 2005-02-10

PRIOR APPLICATION UNMBER: 60/543,296

PRIOR FILING DATE: 2004-02-11

PRIOR FILING DATE: 2004-06-18

PRIOR FILING DATE: 2004-06-18

PRIOR APPLICATION NUMBER: 10/293,418

PRIOR APPLICATION NUMBER: 10/293,418

PRIOR FILING DATE: 2002-11-14

PRIOR APPLICATION NUMBER: 60/331,469

PRIOR APPLICATION NUMBER: 60/340,817

PRIOR APPLICATION NUMBER: 60/340,817
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US-11-054-515-2214
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Sequence 2215, Application US/11054515
Publication No. US2005025532A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Annibodies that Immunospecifically Bind BLyS
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Best Local Similarity
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GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-10-17
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TYPE: PRT
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                                                                                           APPLICATION NUMBER: 60/340,817
FILING DATE: 2001-12-19
APPLICATION NUMBER: 09/880,748
FILING DATE: 2001-06-15
                                           APPLICATION NUMBER: 60/293,499
FILING DATE: 2001-05-25
APPLICATION NUMBER: 60/277,379 FILING DATE: 2001-03-21
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FILING DATE: 2001-03-16
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66.7%;
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Pred. No. 23;
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RESULT 14 US-11-054-515-2219

Sequence 2219, Application US/11054515 Publication No. US2005025532A1 GENERAL INFORMATION:

APPLICANT: Ruben et al

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FILE REFERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR PILING DATE: 2004-02-11
PRIOR PPLICATION NUMBER: 60/580,347
PRIOR PILING DATE: 2004-06-18
PRIOR PILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-054-515-2217
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US-11-054-515-2217
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Best Local S
Matches 4
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Remaining Prior Application data removed - See File Wrapper or NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2217
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PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2215
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Publication No. US20050255532A1
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Best Local Similarity
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FILING DATE: 2001-06-15
APPLICATION NUMBER: 60/293,499
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9 YVĻHYY 14
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66.7%;
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Pred. No. 23;
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RESULT 15
US-11-054-515-2298
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US-11-054-515-2219
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PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
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PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR APPLICATION NUMBER: 60/880,748
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APPLICANT: Ruben et al
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PRIOR FILING DATE: 2000-10-17
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TYPE: PRT
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OR APPLICATION NUMBER: 60/340,817
OR FILING DATE: 2001-12-19
OR APPLICATION NUMBER: 09/880,748
OR FILING DATE: 2001-06-15
OR APPLICATION NUMBER: 60/293,499
OR FILING DATE: 2001-05-25
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OR APPLICATION NUMBER: 60/293,499
OR FILING DATE: 2001-05-25
OR APPLICATION NUMBER: 60/277,379
OR FILING DATE: 2001-03-21
OR APPLICATION NUMBER: 60/276,248
OR FILING DATE: 2001-03-16
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                                        APPLICATION NUMBER: 60/240,816 FILING DATE: 2000-10-17
                                                                                        APPLICATION NUMBER: 60/277,379
FILING DATE: 2001-03-21
APPLICATION NUMBER: 60/276,248
FILING DATE: 2001-03-16
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Prior Application data removed - SEQ ID NOS: 3247
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Pred. No. 23;
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                     File Wrapper or PALM
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Query Match
Best Local Similarity
Matches 4; Conserve
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2002-11-14
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                                                                                                                   Sequence 489, Application US/11033039 Publication No. US20060002947A1 GENERAL INFORMATION:
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Publication No. US20050255532A1
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LENGTH: 16
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PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 09/880,748
APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN
TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01
CURRENT APPLICATION NUMBER: US/11/033,039
CURRENT FILING DATE: 2005-01-11
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PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
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FILING DATE: 2001-05-25
APPLICATION NUMBER: 60/277,379
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    Mismatches

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Pred. No. 23;
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Pred. No. 23;
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RESULT 19
US-10-939-890-11
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US-11-033-039-489
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PRIOR FILING DATE: 1999-09-14
NUMBER OF SEQ ID NOS: 1452
SOFTWARE: PatentIn version 3.3
SEQ ID NO 512
                                                                                   GENERAL
                                                                                                  Sequence 11, Application US/10939890 Publication No. US20050250700A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 512, Application US/11033039 Publication No. US20060002947A1 GENERAL INFORMATION:
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Best Local
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                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN
TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01
CURRENT APPLICATION NUMBER: US/11/033,039
CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 10/245,871
PRIOR APPLICATION NUMBER: 10/245,871
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-09-17
PRIOR PELING DATE: 2002-07-17
PRIOR FILING DATE: 2002-07-17
PRIOR FILING DATE: 2002-07-17
PRIOR FILING DATE: 2002-07-17
PRIOR PRIOR PRIOR DATE: 2002-07-17
PRIOR PRIOR PRIOR DATE: 2002-07-17
                                      APPLICANT: Sato, Aaron K. APPLICANT: Sexton, Danie
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PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: 09/396,813
PRIOR FILING DATE: 1999-09/34
NUMBER OF SEQ ID NOS: 1452
                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: MOD_RES
LOCATION: (5)..(5)
OTHER INFORMATION: Ava
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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TYPE: PRT
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                                                                                 INFORMATION:
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5; Conserva
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                                                                                                                                                                                                                                                                                                        h 25.4%;
Similarity 55.6%;
5; Conservative
Sexton, Daniel J.
Dransfield, Daniel T.
Ladner, Robert C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25.4%;
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Pred. No. 5.8e
2; Mismatches
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Pred. No. 29;
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; CRGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: KDR or VEGF/KDR Binding Polypeptide
US-10-939-890-11
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Best Local S
Matches 5
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NUMBER OF SEQ ID NOS: 883
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/939,890
CURRENT FILING DATE: 2004-09-13
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APPLICANT:
APPLICANT:
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PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: US 10/382,082
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: PCT/US03/06731
PRIOR FILING DATE: 2003-03-03
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PRIOR FILING DATE: 2003-01-15
PRIOR APPLICATION NUMBER: US 60/360,851
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T: Song, Bo
T: Swenson, Rolf E.
T: Von Wronski, Mathew A.
INVENTION: KDR AND VEGF/KDR BINDING PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                  3 HCYLHNG
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Fan, Sudha
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Dransfield, Daniel T.
Ladner, Robert C.
                                                             Shrivastava, Ajay
                                                                                                                                       Khurana, Sudha
Linder, Karen E.
Marinelli, Edmund R.
Nanjappan, Palaniappa
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Shrivastava, Ajay
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Bussat, Philippe
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Pochon, Sibylle
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Pochon, Sibylle
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Marinelli, Edmund R.
Nanjappan, Palaniappa
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b. US20050250700A1
                                                                               Ramalingam, Kondareddiar
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Sudha
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FILE REFERENCE: D0617.70014US00

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PRIOR FILING DATE: 2003-09-11
PRIOR PPLICATION NUMBER: US 10/382,082
PRIOR FILING DATE: 2003-03-03
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: PCT/US03/06731
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US 60/440,411
PRIOR APPLICATION NUMBER: US 60/440,411
PRIOR PILING DATE: 2003-01-15
PRIOR PILING DATE: 2003-03-01
PRIOR FILING DATE: 2002-03-01
                                                                       Query Match
Best Local S
Matches 5
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NUMBER OF SEQ ID NOS: 1452
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 513
LENGTH: 17
TYPE: PRT
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PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: 09/396,813
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CURRENT FILING DATE: 2005-01-11
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: XU, MINZHEN
TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 10/245,871
                                                                                                                                                          NAME/KEY: MOD_RES
LOCATION: (5)...(5)
OTHER INFORMATION: Ava
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: hybrid peptide
                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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5; Conserv
                                    MHYYLNNGA 21
MHYYVSMDA 14
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o. US20060002947A1
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                                                                       Score 30; DB Pred. No. 35; 2; Mismatches
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Pred. No.
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Query Match
Best Local Similarity
Watches 5; Conserve
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US-11-054-515-2811
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                                                                                                                                                                                             FILE REFERENCE: PZ013P2C1
CURRENT APPLICATION NUMBER: US/10/986,501
CURRENT FILING DATE: 2004-11-12
PRIOR APPLICATION NUMBER: US/10/621,363
PRIOR FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/969,730
PRIOR APPLICATION NUMBER: 09/974,639
PRIOR APPLICATION NUMBER: 09/774,639
PRIOR APPLICATION NUMBER: 09/774,639
PRIOR APPLICATION NUMBER: 60/238,291
PRIOR APPLICATION NUMBER: 60/238,291
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PRIOR FILLING DATE: 2004-02-11
PRIOR PELLOATION NUMBER: 60/580,347
PRIOR PELLOATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILLING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILLING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILLING DATE: 2001-12-19
PRIOR FILLING DATE: 2001-12-19
PRIOR FILLING DATE: 2001-12-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 203, Application US/10986501
Publication No. US20050244845A1
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 2000-10-17
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CURRENT FILING DATE: 2005-02-10
                                                                  PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: 09/244,112
PRIOR FILING DATE: 1999-02-04
PRIOR APPLICATION NUMBER: PCT/US98/10
PRIOR FILING DATE: 1998-08-04
PRIOR FILING DATE: 1998-08-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ruben et al.
TITLE OF INVENTION: 90 Human Secreted Proteins
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PRIOR FILING DATE: 2001-06-15
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                                                                        APPLICATION NUMBER: PCT/US98/16235 FILING DATE: 1998-08-04
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                  APPLICATION NUMBER: 60/056,371 FILING DATE: 1997-08-19
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APPLICATION NUMBER: 60/056,732
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; OTHER INFORMATION: Potential Epitope of human Factor VIII
US-10-511-559-254
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 254
                                                                                                                                                                              Sequence 2258, Application US/11054515
Publication No. US20050255532A1
GENERAL INFORMATION:
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SOFTWARB: PatentIn Ver. 2.0
SEQ ID NO 203
LENGTH: 8
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Matches 5; Conservative
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                               APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/511,559
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: PCT/EP03/04063
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: EP 02008712.8
PRIOR PILING DATE: 2002-04-18
PRIOR PILING DATE: 2002-04-18
PRIOR PILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-24
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APPLICANT: CARR, Francis, J.
TITLE OF INVENTION: MODIFIED FACTOR VIII
FILE REFERENCE: MER-133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: JONES, Tim
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APPLICATION NUMBER: 60/580,347 FILING DATE: 2004-06-18
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Search completed: January 26, 2006, 08:39:06 Job time : 4.36207 secs

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NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2258
LENGTH: 16
TYPE: PRT
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Matches 4; Conserv
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PRIOR FILLING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILLING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILLING DATE: 2000-10-17
PRIOR FILLING DATE: 2000-10-17
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PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
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                                         11 YTMHYY 16
9 YVFHYY 14
                                                                                Conservative
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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hypothetical 1.7K
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38K protein 3129 epoxide hydrolase
T-cell receptor de
F7-1 fimbrial prot
19 mu chain V regi
hsp90 protein homo
lambda 112 protein
protein-tyrosine k
celF protein beta c
Ig heavy chain CRD
s-allele-associate
T-cell antigen rec
TSH protein beta c
Ig H chain V-D-J r
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Ig H chain V-D-J r
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A; Cross-references: UNIPARC: UPI00000003A9; GB: V01278; GB: J01755; CA: CROSS-references: UNIPARC: UPI00000003A9; GB: V01278; GB: J01755; CA: CROSS-RESEARCH SERVICES: AREA CONTROL MECHANISM OF The A; Mote: this peptide is involved in the control mechanism of the R; Catchpole, I.; Thomas, C.; Davies, A.; Dyke, K.G.H.

R; Can. Microbiol. 134, 697-799, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Staphylococcus aureus
C;Species: 29-Jul-1981 #sequence revision 01-Sep-1981 #text_change 09-Jul-2004
C;Accession: B93717; B93866; A6568; S03586; A03599
R;Gryczan, T.J.; Grandi, G.; Hahn, J.; Grandi, R.; Dubnau, D.
Nucleic Acide Res. 8, 6081-6097, 1980
A;Title: Conformational alteration of mRNA structure and the posttranscriptional A;Reference number: A93717; MUID:81124320; PMID:6162157
A;Accession: B93717
                                                                                                                                                                                                                                                                                                                        R;Horinouchi, S.; Weisblum, B.

Proc. Natl. Acad. Sci. U.S.A. 77, 7079-7083, 1980
A;Title: Posttranscriptional modification of mRNA conformation: mechanism that regulates A;Reference number: A93866; MUID:81175093; PMID:6938954
A;Accession: B93866
A;Reference number: A46568; MUID:89036120; PMID:3141573
A;Accession: A46568
A;Status: preliminary
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A;Title: Nucleotide sequence of the sex pheromone inhibitor
A;Reference number: A37391; MUID:91261999; PMID:2128961
A;Accession: B37391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: UNIPROT:Q52135; UNIPARC:UPI00000B26AA
A;Note: sequence extracted from NCBI backbone (NCBIN:99901, NCBIP:99906)
R;Clewell, D.B.; Pontius, L.T.; An, F.Y.; Ike, Y.; Suzuki, A.; Nakayama,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    J. Bacteriol. 174, 3152-3160, 1992
A;Title: Conjugative transfer of Enterococcus faecalis plasmid A;Reference number: A41868; MUID:92250408; PMID:1315730
A;Contents: plasmid pAD1
A;Accession: B41868
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C;Accession: B41868; B37391_
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A; Residues: 1-15 < CLE>
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:Residues: 1-19 <GRY>
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Pred. No. 95;
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                                                                                                                                                                                           synthesis of the
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A;Genome: chloroplast C;Superfamily: NADH d C;Keywords: chloropla
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R;Hayashida, N.; Matsubayashi, T.; Shinozaki, K.; Sugiura, M.; Inoue, K.; Hiyama, Curr. Genet. 12, 247-250, 1987
A;Title: The gene for the 9 kd polypeptide, a possible apoprotein for the iron-su A;Reference number: S07170; MUID:88210537; PMID:3329576
A;Accession: S08590
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                                                                                                                                                                                                                                                                                                                                                           NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 4 - Debney's tobacco chloroplast C;Species: chloroplast Nicotiana debney' (Debney's tobacco) C;Date: 25-Feb-1994 #sequence_revision 01-Dec-1995 #text_change 03-Jun-2002 C;Date: 25-Feb-1990 #sequence_revision 01-Dec-1995 #text_change 03-Jun-2002 C;Accession: S08590
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                                                                                                                                                                             A;Molecule type: DNA
A;Residues: 1-21 <HAY>
A;Cross_references: UNIPARC:UPI0000000AD9; EMBL:X05881; NID:g11790; PIDN:CAA29305.1;
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A;Accession: B26930
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A; Residues: 1-19 < MAY>
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A;Residues: 1-19 <MON>
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J. Bacteriol. 169, 340-350, 1987
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;Species: Bacillus sphaericus
;Date: 08-Mar-1989 #sequence_revision 08-M
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Best Local
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4; Conserv
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milarity 33.3%;
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nitrogenase cofactor synthesis protein nifs - Anabaena variabilis (fragment) N;Contains: L-cysteine sulfurtransferase (EC 2.8.1.-) C;Species: Anabaena variabilis C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004 C;Accession: S32677 R;Monnerjahn, U.; Boehme, H. submitted to the EMBL Data Library, December 1992 submitted to the EMBL Data Library, December 1992
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-15 <MON:
                                                              A;Description: Cloning and A;Reference number: S32675 A;Accession: S32677
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A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0254
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A;Residues: 1-20 <CHI:>
A;Residues: 1-20 <CHI:>
A;Cross-references: UNIPROT:Q7M447; UNIPARC:UPI00001754DB
C;Superfamily: glutathione transferase
C;Keywords: eye lens
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J. Protein Chem. 7, 527-534, 1988
A;Title: The protein sequence homology of gamma-crystallins among major vertebrate class A;Reference number: A60894; MUID:89351593; PMID:3255376
A;Accession: E60894
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A; Residues: 1-14 < YAM>
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C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0254
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C;Species: Sepia esculenta (Pacific cuttlefish)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1
C;Accession: E60894
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cross-references: UNIPARC:UPI000017C1EB
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22.9%;
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                                                                                                       Library, December 1992 expression in E. coli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunoglobulin
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Pred. No. 3.8e+02;
2; Mismatches 3;
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Pred. No. 2.7e+02;
3; Mismatches 3;
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                                                                                                          Anabaena
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RESULT 10
S63503
C;Species: Eubacterium acidaminophilum
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
C;Accession: S63503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig H chain V-D-J region (clone B-less 151) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                                   A; Experimental source: C; Keywords: immunoglob
                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPARC:UPI000017C694
A;Experimental source: bone marrow pre-B
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: DNA
A: Residues: 1-14 < LEV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Accession: G64003

R;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.; Kerlavage, Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; Weidman, D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen, N.S.M. Science 269, 496-512, 1995
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                                                                        83K protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Accession: PH1628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mic A;Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Levinson, D.A.; Campos-Torres, J.; Leder, J. Exp. Med. 178, 317-329, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd. A;Reference number: A64000; MUID:95350630; PMID:7542800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein HI0195 - Haemophilus influenzae (strain Rd KW20)
C;Species: Haemophilus influenzae
C;Datc: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 30-Jun-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Cross-references: UNIPARC: UPI000017AAED;
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A;Residues: 1-12 <TIGR>
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                                                                        Eubacterium acidaminophilum (fragment)
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57.1%;
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Pred. No. 5.50
0; Mismatches
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Pred. No.
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Pred. No. 4.1e+02;
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5.5e+02;
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C;Accession: S42282
R;Crickmore, N.; Wheeler, V.C.; Ellar, D.J.
Mol. Gen. Genet. 242, 365-368, 1994
Mol. Gen. Genet. 242, 365-368, 1994
A;Title: Use of an operon fusion to induce expression and crystallisation of a Bacillus A;Reference number: S42282; MUID:94150472; PMID:7906381
A;Accession: S42282
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A;Title: Molecular cloning and expression of the transformation sensitive epithelial ma A;Reference number: S34753; MUID:93294871; PMID:8515476
A;Accession: S37684
                                                         Query Match
Best Local Similarity
Marches 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein IEF SSP 9125 - human (fragments)
C;Species: Homo sapiens (man)
C;Date: 09-Dec-1993 #sequence_revision 17-Nov-1995 #text_change 17-Nov-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Meyer, M.; Granderath, K.; Andreesen, J.R. Eur. J. Biochem. 234, 184-191, 1995
A;Title: Purification and characterization of protein P(B)
В
                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Bacillus thuringiensis
C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997 #text_change
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A;Molecule type: protein
A;Residues: 1-20 <LEF>
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A; Residues: 1-17 <MEY>
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                                                                                                                                                                                                                            A;Cross-references: UNIPARC:UPI000017ACD4
A;Experimental source: subsp. galleriae 916
                                                                                                                                                                                                                                                                  A; Molecule type: protein
A; Residues: 1-10 < CRI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                         parasporal crystal protein cryIIB - Bacillus thuringiensis plasmid (fragment)
N;Alternate names: delta-endotoxin
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                                                                                                                                                   ;Genome: plasmid
;Keywords: delta
                                                                                                                                                                                       Gene: cryIIB
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Local Similarity 44.4%;
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     MNTVLNNG
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Pred. No. 9.7e+02;
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Pred. No. 6.8e+02;
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Pred. No. 5.
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5.5e+02;
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D-arabinose 1-dehydrogenase [NAD(P)] (EC 1.1.1.117) - yeast (Candida albicans) C;Species: Candida albicans albicans C;Species: Candida albicans (C;Date: 29-Jan-1998 #sequence_revision 06-Feb-1998 #text_change 03-Jun-2002 C;Accession: S72217 R;Kim, S.T.; Huh, W.K.; Kim, J.Y.; Hwang, S.W.; Kang, S.O. Biochim. Biophys. Acta 1297, 1-8, 1996 A;Title: D-arabinose dehydrogenase and biosynthesis of erythroascorbic acid in A;Reference number: S72217; MUID:96439039; PMID:8841374 A;Accession: S72217
                                                                                A;Molecule type: protein
A;Residues: 1-14 <KIM>
A;Cross-references: UNIPARC:UPI000017CDB9
C;Keywords: oxidoreductase
                                                                                                                                                                                                                                                                                                                                                            RESULT 15
S72217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPARC:UPI000017C69E
A;Experimental source: bone marrow pre-B lymphocyte
C;Keywords: immunoglobulin
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A;Title: Molecular characterization of transgene-induced A;Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice
A;Reference number: PH1580; MUID:93301609; PMID:8315387
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C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
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A; Molecule type: DNA
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A; Residues: 1-16 < LEV >
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les 3; Conserv
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                   20.3%;
Score 24; DB 2;
Pred. No. 1.1e+03;
1; Mismatches 1
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Pred. No. 9.1e
2; Mismatches
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Pred. No.
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Ripavison, M.D.; Rixon, F.J.; Davison, A.J.
J. Gen. Virol. 73, 2709-2713, 1992
A;Title: Identification of genes encoding two capsid proteins A;Reference number: PQ0544; MUID:93019027; PMID:1328483
A;Accession: PQ0544
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J. Exp. Med. 173, 395-407, 1991
A:Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A:Reference number: PT0222; MUID:91108337; PMID:1899102
A:Accession: PT0332
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A;Title: Structural and immunological studies on the soluble formate dehydrogenase from A;Reference number: S59492; MUID:96145736; PMID:8561915
A;Accession: S59495
                                       A; Molecule type: protein A; Residues: 1-20 < DAV>
                                                                                                                                                                                     capsid protein VP5 - human herpesvirus 1 (fragments)
C;Species: human herpesvirus 1
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 07-May-1999
C;Accession: PQ0544
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C;Keywords: heterotetramer;
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A; Residues: 1-19 < YAM>
A; Cross-references: UNIPARC: UPI000017C21A
A; Experimental source: B lymphocyte
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
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PT0332
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S59495
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Date: 27-Apr-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
  ;Cross-references: UNIPARC:UPI000017A7CA;Experimental source: strain 17
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Experimental source: strain
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Best Local
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1es 3; Conservation
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Pred. No. 1.6e+03;
"'amatches 2;
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Pred. No. 1.1e+03;
0; Mismatches 1
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RESULT
PH1607
                                                                Ig H chain V-D-J region (wild-type clone 333) - mouse (fragment) (Species: Mus musculius (house mouse) C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change C;Accession: PH1607
R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mic-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular haracterization of transgene-induced immunodeficiency in B-less mic A;Reference number: PH1580; MUID:93301609; PMID:8315387
A;Accession: PH1601
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PH1601
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A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobuli
                                                                                                                                                                                                                                                                                                                                                                                                             C; Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Experimental source:
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A; Residues: 1-14 <LEV>
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C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994
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A;Title: Preferential utilization of specific immunoglobulin he A;Reference number: PT0222; MUID:91108337; PMID:1899102
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C;Keywords: capsid protein
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A;Residues: 1-13 <YAM>
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;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993
;Accession: PT0263
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Best Local Similarity
Matches 4; Conserv
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Local Similarity 50.0%;
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Pred. No. 1.6e+03;
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Pred. No. 1.5e+03;
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Pred. No. 1.7e+03;
1; Mismatches 0
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                                                                                    #text_change 17-Mar-1999
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RESULT 24
PQ0751
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J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1307
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B24735
                                                                                                                                                                                                                                                     A;Cross-references: UNIPARC:UPI0000176935
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Accession: B24735

R;Mannervik, B.; Alin, P.; Guthenberg, C.; Jensson, H.; Tahir, M.K.; Warholm, Proc. Natl. Acad. Sci. U.S.A. 82, 7202-7206, 1985

A;Title: Identification of three classes of cytosolic glutathione transferase A;Reference number: A24735; MUID:86042634; PMID:3864155

A;Accession: B24735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 glutathione transferase (EC 2.5.1.18) 1-2 - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Oct-1987 #sequence_revision 25-Oct-1987 #text_change 18-Jun-1993
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A;Experimental source: bone marrow pre-B lymphocyte
C;Superfamily: immunoglobulin V region; immunoglobulin homology
self-incompatibility protein S4 -
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A; Residues: 1-19 <WAS>
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PH1307
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A;Residues: 1-18 <MAN>
A;Cross-references: UNIPARC:UPI000017542B
C;Superfamily: glutathione transferase
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A; Residues: 1-17 <LEV>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig heavy chain DJ region (clone C96-119) - human (fragment)
                                                                                                                                                                            Query Match
Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                ;Species: Homo sapiens (man)
;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999;Accession: PHI307
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3; Conserv
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Similarity 50.0%;
4; Conservative
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Pred. No. 2e+(
0; Mismatches
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Pred. No. 2.3e+03;
2; Mismatches 1
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potato (fragment)
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A;Genome: chloroplast
C;Superfamily: H(+)-transporting ATP synthase protein 6
C;Keywords: ATP biosynthesis; chloroplast; hydrolase; membrane-associated complex; thyla)
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FEBS Lett. 377, 163-166, 1995
A;Title: Isolation of CF(0)CF(1) from Chlamydomonas reinhardtii cw15 and the N-terminal &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         H+-transporting two-sector ATPase (EC 3.6.3.14) chain I - Chlamydomonas reinhardtii chlon
N;Alternate names: ATP synthase chain I
C;Species: chloroplast Chlamydomonas reinhardtii
C;Date: 04-Dec-1997 #sequence_revision 12-Dec-1997 #text_change 31-Dec-2004
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S68392
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C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 31-Dec-2004
C;Accession: PQ0751
Search completed: January 26, 2006, 08:05:02 Job time: 12.5 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Kirch, H.H.; Uhrig, H.; Lottspeich, F.; Salamini, Theor. Appl. Genet. 78, 581-588, 1989
A;Title: Characterization of proteins associated will
                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPARC:UPI0000175ED3
A;Experimental source: strain CW15
                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: protein A; Residues: 1-11 <FIE>
                                                                                                                                                                                                                                                                                                                                                                                                                                A;Reference number: $68388; MUID:96128220; PMID:8543042 A;Accession: $68392
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A;Experimental source: style
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A; Accession: PQ0751
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MEDLINE=21621309; PubMed=11751136; DOI=10.1128/AAC.46.1.211-215.2002; Matsuoka M., Inoue M., Nakajima Y., Endo Y.;

MINEW erm Gene in Staphylococcus aureus clinical isolates.";

NAtimicrob. Agents Chemother. 46:211-215(2002).

EMBL; ABO14481; BAB20747-1; -; Genomic DNA.

SEQUENCE 19 AA; 2195 MW; 605CBAD99E4B6B7F CRC64;
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Leader peptide of ErmGM.
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"The staphylococcal tetracycline macrolide-
resistance plasmid pSTE2 is an RSA mediated
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13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
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body boundary proteins (oleosins) from the dev
sunflower (Helianthus annuus L.).";
Biochem. J. 314:333-337(1996).
SEQUENCE 15 AA; 1619 MW; CDC28A062F7F8704
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EMBL; CAAJ01003410; (
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Eukaryota; Alveolata;
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ORFNames=PC106693.00.0;
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MEDLINE=96195148; PubMed=8660304;
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Science 307:82-86 (2005).

Science 307:82-86 (2005).

Science 307:82-86 (2005).
                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.

MEDLINE=92250408; PubMed=1315730;

Pontius L.T., Clewell D.B.;

Pontius L.T., Clewell D.B.;

"Conjugative transfer of Enterococcus faecalis
nucleotide sequence and transcriptional fusion
involved in positive regulation.";

J. Bacteriol. 174:3152-3160(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE.

MEDLINE-91261999; PubMed=2128961;

MEDLINE-91261999; PubMed=2128961;

Clewell D.B., Pontius L.T., An F.Y., Ike Y., Suzuki

"Nucleotide sequence of the sex pheromone inhibitor
of Enterococcus faccalis conjugative plasmid pAD1.";
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EMBL; M62888; AAA98040.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plasmid pAD1.
Plasmid pAD1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORF; putative.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=5821;
                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=2520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
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Eukaryota; Alveolata;
                                                                                                                                                                                                                                               PIR; B41868; B41868.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORFNames=PB105101.00.0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAAI01002093; CAH97654.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         _9ZZZZ PRELIMINARY;
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                                                                                                                          Similarity
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  B IXAHALAW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NVMXYVLYTMYYLVNN 19
                                            TAXHWLAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SILIYAIYT -- FYFNH 16
                                                                                                                                                                                                   15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 AA; 2419 MW;
                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                   $
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plasmids.
                                                                                                                                                                                                   1874 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27.1%;
                                                                                                                     26.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Last sequence up
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Created)
                                                                                                Score 31; DB 2;
Pred. No. 7.8e+0;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 32; DB 2; Le
Pred. No. 7.3e+02;
6; Mismatches 3;
                                                                                                                                                                                                                                                                      Genomic_DNA
                                                                                                                                                                                                   0D9D07E3079E3559 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3ABE7624B6EF6EA8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -; Genomic_DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence update)
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                                                                                                                          8e+02;
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analysis of a region
                                                                                                                                               Length 15
                                                                                                  Indels
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(iAD1)
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D1) determinant
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RESULT
Q4Z5L3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  James K., Rutherford K., Harris B., Harris D., Christophides G.K.,
RA James K., Rutherford K., Harris B., Harris D., Churcher C.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
RA Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.C.,
RA Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.,
T' A comprehensive survey of the Plasmodium life cycle by genomic,
transcriptomic, and proteomic analyses.";
L Science 307:82-86(2005).
C --- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
preliminary data.
Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 4
                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
Hall N. Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Berriman M., Florens L., Janssen C.S., Pain A., Christophides G.
James K., Rutherford K., Harris B., Harris D., Churcher C.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos
Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.;
"A comprehensive survey of the Plasmodium life cycle by genomic,"
                                                                                                                                                                                                                                                                                                                                             13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q4YVN6;
                                                                                                                              transcriptomic, and protec
Science 307:82-86(2005)
-i- CAUTION: The sequence
                                                                                                                                                                                                                                                                                               Plasmodium berghei.
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
                                                                                                                                                                                                                                                                                                                                                                                                    Q4Z5L3_PLABE PRELIMINARY;
Q4Z5L3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         preliminary data.
EMBL; CAA101002238; CAH97920.1; -;
Hypothetical protein.
SEQUENCE 21 AA; 2441 MW; B79B0
                                                               Hypothetical protein NON_TER 1
                                                                                           EMBL;
                                                                                                                                                                                                                                                                                                                                    ORFNames=PB101034.00.0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical protein. ORFNames=PB105419.00.
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                                                                                                                                                                                                                                                                                         NCBI_TaxID=5821;
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13-SEP-2005
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                                                                                          preliminary data.
; CAAI01000433; CAH94418.1;
                                                                                                                    EMBL/GenBank/DDBJ whole
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Similarity 6; Conserv
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                                                   A
A
                                                                                                                                                           and proteomic analyses.";
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                                                    2302
           25.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31, Createu,
31, Last sequence update)
31, Last annotation update)
                                                    Μ¥.
                                                                                                                                  shown
                                                                                                                   genome
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           Score 30.5; DB 2;
Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B79B04782E2A5804 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31;
                                                    28C71708F618BA20 CRC64;
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                                                                                                                              here
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                                                                                        Genomic_DNA
                                                                                                                   shotgun
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                                                                                                                                  is derived
                                                                                                                                                                                                                                                                                                                                                                                                                   18
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                          DB 2;
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                                                                                                                      (WGS)
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                                                                                                                                  from
                                                                                                                                                                                                                                     Christophides G.K.,
                        Length
                                                                                                                      entry which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                           18;
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Conservative

4.

Mismatches

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ID4XG09
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RRRRRRRRRRRRRROOGGEEDTTE
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Best Local S
Matches 5
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Plasmodium chabaudi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUL-1986
21-JUL-1986
10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LPRM STAAU
P03063;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A., Berriman M., Florens L., Janssen C.S., Pain A., Christophides G.K., James K., Rutherford K., Harris B., Harris D., Churcher C., Janes R., Rutherford K., Doggett J., Trueman H.E., Mendoza J., Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J., Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.C. Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.; "A comprehensive survey of the Plasmodium life cycle by genomic,"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q4XG09
                                                                                                                                                                                       MEDLINE=81124320; PubMed=6162157; Gryczan T.J., Grandi G., Hahn J., Grandi R., "Conformational alteration of mRNA structure
                                                                                                                                                                                                                                                                                                                                                                            Staphylococcus aureus.
Plasmid pE194, and Plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STAAU
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Hypothetical protein.

15 AA; 1776 MW;
                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
PLASMID=pE194;
                                                                                                                                                                                                                                                                                                                                                                                                                       Name=ermC;
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                                                  PLASMID=pE194
                                                                                                                                                               posttranscriptional regulation of erythromycin-induced drug
                                                                                                                                                                                                                                                                                                                                                           Bacteria; Firmicutes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
                            MEDLINE=81175093; PubMed=6938954;
                                                                          NUCLEOTIDE SEQUENCE
                                                                                                                      Nucleic
                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=1280;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide)
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; CAAJ01006563; (
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                                                                                                                        Acids
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       s;
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(Rel. 01, Last sequence update)
(Rel. 47, Last annotation update)
chylase leader peptide (Erythromyc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                      Res.
       Weisblum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
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                                                                                                                      8:6081-6097(1980)
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83.3%;
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Bacillales;
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Pred. No. 1.1e+03;
1; Mismatches 0
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EMBL; M3784; AAA98225.1; -; Genomic_DNA.
EMBL; M19652; AAA20193.1; -; Unassigned_D
PIR; B93717; LFSAP9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=86037248; PubMed=2414456; Mayford M., Weisblum B.; "Messenger RNA from Staphylococcus aureus that specifies "Messenger RNA from Staphylococcus aureus that specifies lincosamide-streptogramin resistance. Demonstration of it conformations and of the leader peptide it encodes."; J. Mol. Biol. 185:769-780(1985).
NUCLEOTIDE SEQUENCE.
MEDLINE=96379895; PubMed=8787908;
Lodder G., Schwarz S., Gregory P., Dyke K.
"Tandem duplication in ermC translational
                                                           Bacteria; Firmicutes;
NCBI_TaxID=29383;
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Plasmid pSES6
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RESULT 13
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Submitted (MAY-1998) to the ENEMBL; Y17294; CAB46339.1; -; GHypothetical protein; Plasmid. SEQUENCE 19 AA; 2237 Mu.
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EMBL; X82668; CAA57984.1
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Bacteria; Firmicutes;
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EMBL; AJ313523; CAC66409.1; -;
Hyporhetical protein; Plasmid.
SEQUENCE 19 AA; 2237 MW; 1
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DOI=10.1126/AAC.46.11.3621-3623.2002;
Schwarz S., Kehrenberg C., Ojo K.K.;
"Staphylococcus sciuri gene erm(33), encoding inducible remacrolides, lincosamides, and streptcyramin B antibiotics, product of recombination between erm(C) and erm(A).";
product of recombination between erm(C) and erm(A).";
Antimicrob. Agents Chemother. 46:3621-3623(2002).
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MEDLINE=87083389; Po
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"Cloning and analysis of ermG, a new mac streptogramin B resistance element from J. Bacteriol. 169:340-350(1987).

EMBL; M15332; AAA22418.1; -; Genomic_DNF pIR; B26930; B26930.
                                                                                                                                                                                                                                                                                                               Submitted (NOV-1994) to the | EMBL; Y09001; CAA70205.1; -; EMBL; X82667; CAA57982.1; -;
                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                    Schwarz
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lodder G., Schwarz S., Gregory P., Dyke K.;
"Tandem duplication in ermC translational attenuator
lincosamide-streptogramin B resistance plasmid pSES6
Staphylococcus equorum.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
MEDLINE=96379895; PubMed=8787908;
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STRAIN=736153; PLASMID=pSES5;
Schwarz S.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lodder G., Werckenthin C., Schwarz S., Dyke K.; "Molecular analysis of naturally occurring ermC-encoc staphylococci isolated from animals with and without with macrolide / lincosamide antibiotics.";
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GN Name-erm
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Q79DT6;
Q79DT6;
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05-JUL-2004 (TrEMBLrel. 27, Last annot Leader peptide.
Name=ermGT;
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Plasmid pRJ5.
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SEQUENCE
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J. Gen. Microbiol. 139:1461-1467(1993).
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"The complete sequence of plasmid pWBG738 from the ST1-MRSA-IV lineage of community methicillin-resistant Staphylococcus aureus."; Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.
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RESULT 21
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DQ4Y2F8_PLACH PRELIMINARY;
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NCBI_TaxID=28406;
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NUCLEOTIDE SEQUENCE.

MEDLINE-94224982; PubMed-8171126; DOI=10.1006/plas.1994.1007;

Tannock G.W., Luchansky J.B., Miller L.A., Connell H.,

Thode-Andersen S., Mercer A.A., Klaenhammer T.R.;

"Molecular characterization of a plasmid-borne (pGT633) erythromycin resistance determinant (ermGT) from Lactobacillus reuteri 100-63.";

Plasmid 31:60-71(1994).

EMBL; M64090; AAA98095.1; -; Genomic_DNA.
                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                     Hypothetical NON_TER
                                                                                                                                                                                                               EMBL;
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Name=PY06333; Synonyms=PY01707;
Plasmodium yoelii yoelii.
Eukaryota; Alveolata; Apicomplex
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01-MAR-2004 (TrEMBLrel 26, Last sequence up
01-MAR-2004 (TrEMBLrel 26, Last annotation
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                                                                                                                                                                                                           preliminary data.
; AABL01002128; EAA18531.1; -; Genomic_DNA
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Pred. No. 1.4e+03;
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13-SEP-2005
13-SEP-2005
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Hypothetical protein.
SEQUENCE 21 AA; 26
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                                                                                         SEQUENCE
                                                                                                      Chloroplast.
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                                                                                                                                                              chloroplast
                                                                                                                                                                           iron-sulfur
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                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons; asterids; lamids; Solanales; Solanaceae; Nicotiana.
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Nicotiana tabacum
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Q7GEY8;
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Science 307:82-86(2005).
-!- CAUTION: The sequence shown here is derived
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Berriman M., Florens
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Eukaryota; Alveolata;
NCBI_TaxID=5825;
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ORFNames=PC102749.00.0;
                                                                                                                                                                        "The gene for the 9 kd polypeptide, iron-sulfur centers A and B of the p
                                                                                                                                                                                                  Hiyama
                                                                                                                                                                                                                       STRAIN=Bright yellow 4; TISSUE=Leave; MEDLINE=88210537; PubMed=3329576;
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L; CAAJ01001590; CAH76652.1; -; Genomic_
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X05881; CAA29305.1; -;
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147 9MOLL
Q7M447 9MOLL 1
Q7M447;
Q1-MAR-2004 (7
                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
MEDLINE=99152621; PubMed=10028295;
Dowton M., Austin A.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Hymenoptera; Apocrita; Ichneumonoidea; Braconidae; Rogadinae; Spinaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Cytochrome oxidase II (Fragment).
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Q9ZYB7_9HYME PRELIMINARY;
Q9ZYB7;
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MEDLINE=89351593; PubMed=3255376;
Chiou S.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sepia esculenta.
Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea;
Decapodiformes; Sepioidea; Sepiidae; Sepia.
                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                      Mitochondrion.
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G
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                                                                                                                                                                                                                                                                                                                       Mol. Biol.
                                                                                                                                                                                                                                                                                                                                                                   "Evolutionary dynamics of a mitochondrial rearrangement 'hot spot'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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01-MAR-2004
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Biol. Evol. 16:298-309(1999).
L; AF072122; AAD17665.1; -; Gen
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ein Chem. 7:527-534(1988).
                                           DEXNVMXYVLYTMHYY 16
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ESVDFLMYLNWVMNYY 19
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20 AA;
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(TrEMBLrel. 26, Last sequence update)
(TrEMBLrel. 26, Last annotation update)
                                                                                              Conservative
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Pred. No. 2.2e+03;
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RESULT 25

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AC Q4X688 p
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DT 13-SEP-2
DT 13-SEP-2
DT 13-SEP-2
DE Hypothet
GN ORFNames
OS Plasmodi
OC Eukaryot
OX NCBI Tax
RN [1]
RA Hall N.,
RA Berriman
RA James K.
RA Quail M.
RA Bidwell
RA Janse C.
RT "A compx
RT "A compx
RT L Janse C.
RT CAUBLE
CC PARBL
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Search completed: January 26, Job time: 71.4052 secs
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Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Hall N., Karras M., Raine J.D., Carlton J.M., Christophides G.K.,
Berriman M., Florens L., Janssen C.S., Pain A., Churcher C.,
James K., Rutherford K., Harris B., Harris D., Churcher C.,
Juail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.C.,
Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.;
"A comprehensive survey of the Plasmodium life cycle by genomic,"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QAX6EB, PLACH PRELIMINARY; PRT; 21 AA. QAX6EB; 13-SEP-2005 (TrEMBLrel. 31, Created) 13-SEP-2005 (TrEMBLrel. 31, Last sequence update) 13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical protein (Fragment).
ORFNames=PC405819.00.0;
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Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium
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L; CAAJ01009531; CAH87528.1; -;
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1: geneseqp1980s:*
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Copyright (c) 1993 - 2006 Compugen Ltd
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ADM29132
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ar84471 Hepacitis

ay73124 Hepacitis

ay73124 Hepacitis

aj00049 Hepacitis

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aj01621 Hepacitis

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aj01110 Hepacitis

aj01256 Hepacitis

aj02138 Hepacitis

aj02138 Hepacitis

bj00088 HCV deriv

bc00953 B7-like s

bj00257 Hepacitis

aj0279584 CTL epito

aj00257 Hepacitis

aj01752 Hepacitis

aj01752 Hepacitis
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1 Cytotoxic
0 Hepatitis
7 HLA bindi
7 Tryptic p
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Tryptic p
hTRT MHC
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Human Alz
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Plasma se
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ANY52519
ID ANY5252
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                                            Sequences AAY52510-Y52522 represent proteolytic fragments of CC Dermatophagoides farinae high molecular weight mite allergen protein (HWM CC -map) composition. The HMW-map composition was isolated from a D. farinae CC presence of proteins that bound to IgE present in mite-allergic dog CC antisera. The HMW-map composition comprises mapA (a 109 kD protein) and CC mapB (98 kD). Mite allergenic proteins and peptides, and nucleic acids cencoding them, may be used in therapeutic compositions to modify an CC animal's hypersensitivity reaction to mite allergens. Animals that may be CC treated include mammals and birds, especially felines, canines, equines, other pets, and work or domestic animals. The proteins or CC fragments may also be used to diagnose allergies via a skin test. The CC proteins and peptides can also be used to raise antibodies, which have a CC variety of potential uses. For example, they can be used as vaccines to CC passively immunise animals against dust mite hypersensitivity, as CC positive controls in test kits and as tools to recover desired dust mite allergens from a mixture of proteins
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66
86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel high molecular weight Dermatophagoides nucleic acid polypeptides used to modify an animals' hypersensitivy to mite allergens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY52519 standard; peptide; 17 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HESK-) HESKA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-MAY-1998;
02-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-FEB-2000
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98US-0085295P.
98US-0098909P.
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ABJ05782
AAU95434
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Aau 95434
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2 Streptoco
4 Streptoco
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Query Match Best Local Similarity

97.6%; 100.0%;

Score 81; Pred. No.

DB 3; Length 17, 4.1e-08;

RESULT 3
AAW45138
ID AAW4
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AAW45138 standard; peptide; 16

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Sequence 17 AA;

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RESULT 2
AAU96323
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                                                                                                                                                                                                                                                                                                   Dermatophagoides, designated Der HMW-map protein, and its related nucleic acid. The Der HMW-map protein is useful for eliciting an immune response against Der HMW-map protein. The protein or a reagent comprising a non-proteinaceous epitope is useful for identifying an animal (e.g., dog, cat) susceptible to or having an allergic response to a mite. A therapeutic composition is useful for desensitising a host animal to an allergic response to a mite. The DNA and protein can be used in the detection of anti-Der HMW-map antibodies in animal fluids, and inhibition of immunoglobulin (Ig) E or Der HMW-map protein activity associated with a disease. Antibodies that bind to Der HMW-map are useful for inhibiting binding of proteins to IgE, to prevent immunocomplex formation, thus reducing hypersensitivity responses to mite allergens, and as vaccines against mite allergen hypersensitivity. Sequences AAU96314-AAU96342 represent Der HMW-map polypeptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Der HMW-map; American house dust mite; antiallergic; mite; IgE; mite allergenic protein; immunoglobulin E; hypersensitivity;
                                                                                                                                                                                                                                               Sequence 17 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New mite allergenic protein isolated from Dermatophagoides, designa
Der HMW-map protein, useful as a vaccine for treating mite allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-351888/38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunocomplex formation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 12; Page 71; 161pp; English.
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17; Conserv
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DKLVMGVPFYGRAXSIE 17
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                                                                                                                                                      97.6%; Score 81; 100.0%; Pred. No.
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RESULT 4
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KW HCV;
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                                                                                                                                                                                                                                                                                                           autoantigen in rheumatoid arthritis. The invention relates to peptides consisting of 16-55 amino acid residues comprising at least one of the following 19 sequences: LVCYYTSWS; FLCTHIIYS; IIYSFANIS; LKTLLSVGG; FIKSVPPF1; FDGLDLAWL; LYVEGRROKG; YDIAKISQH; LDFISIMTY; FISIMTYDF; FRGQEDASP; YAVGYMLRL; MLRIGAPAS; LAYYEICDF; LRGATVHRT; YLKORQLAG; LAGAMVMAL; VMALDLDDF; or LDLDDFQGS. They can be used medically in antigen specific immunosuppressive therapy, particularly the treatment of T-cell mediated destruction of articular cartilage in autoimmune diseases (e.g. rheumatoid arthritis). They can also be used to detect activated autoreactive T cells in an individual. The peptides have a specific immunosuppressive drugs, and do not cause toxic side effects. The peptides are predominantly recognised by autoreactive T cells from rheumatoid arthritis patients, but rarely by those from healthy donors
                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide suitable for use in antigen specific immunosuppressive therapy resembles or mimics epitope present on HC gp-39, so inducing systemic immunological tolerance to rheumatoid arthritis auto:antigen.
 HCV; hepatitis C virus; virucide; vaccine; MHC; HLA;
                           Synthetic peptide C127 derived from a conserved region of HCV
                                                           17-JUN-2004
                                                                                        ADL26336;
                                                                                                                ADL26336 standard; peptide; 15
                                                                                                                                                                                                                                                                                                           Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        epitope present on human cartilage glycoprotein autoantigen in rheumatoid arthritis. The inventi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence
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                                                                                                                                                                                                                                                             56.6%;
                                                                                                                                                                                                                                                               Score 47; DB 2
Pred. No. 0.12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a peptide which
                                                                                                                                                                                                                                                   Mismatches
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Best Local
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27-FEB-2003;
11-JUL-2003;
                                                                                                  The invention relates to a novel method for isolating Hepatitis C Virus (HCV) peptides (HPs). The method of the invention has virucide activity, and may be useful in producing a vaccine. The method is useful for isolating Hepatitis C Virus peptides (HPs) which have a binding capacity to a MHC/HLA molecule or a complex comprising the HCV-peptide and the MHC/HLA molecule for preparing a vaccine against HCV infection. The T cell clone or a T cell population or preparation is useful for identifying heteroclitic epitopes or for preparing a composition for treating HCV infection. The present sequence represents a synthetic peptide derived from a conserved region of HCV.
                                                                                                                                                                                                                                                                              Isolating Hepatitis C Virus peptides (HPs) which have a binding capacity to a MHC/HLA molecule or a complex comprising the HCV-peptide and the molecule by separating the complex from the HCV-peptides which do not bind to the molecule.
                                                                                                                                                                                                                                                                                                                                               idm;
                                                                            Sequence 15 AA;
                                                                                                                                                                                                                                                    Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  major histocompatibility complex; human leukocyte antigen
                                                                                                                                                                                                                                                                                                                                                                        Zauner W,
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                                                                                                                                                                                                                                                                                                                                               2004-269899/25
 7 VPFYGRAXSIE 17
                        Similarity 7; Conserv
                                                                                                                                                                                                                                                    1; Page 31;
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                                                                                                                                                                                                                                                                                                                                                                        Zinke S,
                                                                                                                                                                                                                                                                                                                                                                                Habel A,
                           Conservative
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2003WO-EP002005
2003EP-00450171
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                                       50.6%;
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                                       Score 42; D
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RESULT 5
                                                      DNA vaccine; vaccine; antigen gene; antigen; hepatitis C virus; HCV; recombinant adenovirus vaccine; adenovirus; HCV infection; hepatitis C virus infection; infection; virucide.
Hepatitis C
Synthetic.
                                                                                                                                   Hepatitis C
                                                                                                                                                                            01-JUL-2004
                                                                                                                                                                                                                ADM29133;
                                                                                                                                                                                                                                                      ADM29133 standard;
                  virus
                                                                                                                                   virus cellular immune response related
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08-APR-2004.

WO2004028563-A1

24-SEP-2003; 2003WO-KR001951

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RESULT 6
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Best Local
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24-NOV-2000;
28-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a DNA vaccine which comprises a plasmid containing 2-6 kb of the total antigen gene of hepatitis C virus (HCV). Also described: (1) a recombinant adenovirus vaccine including an adenovirus containing 2-6 kb of total antigen gene of HCV; (2) a vaccine administrating method; and (3) a method for treating or preventing HCV infection. The DNA vaccine has virucide activity. The DNA vaccine is useful in treating or preventing HCV infection. The present sequence represents a peptide from the HCV peptide pool used for the investigation of cellular immune response in the exemplification of the present
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06-NOV-2002;
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(DONG-)
(DAEW-)
  Screening, diagnosis or prognosis of vascular dementia (VD), useful
                                                                                           Herath
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                                                                                                                                                                                                                                                                                                                                 20-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                               WO200169261-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis; prognosis; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vascular Dementia; VD; VD-associated protein isoform; VPI; screening;
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                                                                                           HMAC,
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DONG-A PHARM CO LTD

DAEWOONG CO LTD.
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                                                                                                                                         OXFORD GLYCOSCIENCES UK LTD
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                                                                                                                                                                                       2000GB-00006285.
2000GB-00028734.
2000US-00724391.
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2002KR-00068496.
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                                                                                              Parekh
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                                                                                           Rohlff
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 42;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein isoform (VPI)
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     determining stage of VD and monitoring the effect of VD therapy, comprises analyzing body fluid by 2-dimensional electrophoresis for features correlated with VD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to screening, diagnosis or prognosis of Vas Dementia (VD) in a subject comprising analysing body fluid from t subject by 2-dimensional (2-D) electrophoresis to generate a 2-D
                                                                                    Screening, diagnosis or prognosis of vascular dementia (VD), useful determining stage of VD and monitoring the effect of VD therapy, comprises analyzing body fluid by 2-dimensional electrophoresis for features correlated with VD.
                                                                                                                                                                                                                                                  15-MAR-2000; 2000GB-00006285
24-NOV-2000; 2000GB-00028734
28-NOV-2000; 2000US-00724391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vascular dementia-associated protein isoform (VPI) 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABB56114 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 6;
                                                                                                                                                              WPI; 2001-557937/62.
                                                                                                                                                                                                                                                                                                                                          20-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                       WO200169261-A2
                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis; prognosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Vascular Dementia; VD; VD-associated
                                                                                                                                                                                                                                                                                                            14-MAR-2001; 2001WO-GB001106
                                                                                                                                                                                                                        (OXFO-) OXFORD GLYCOSCIENCES UK LTD
                                                                                                                                                                                           HMAC,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page
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LVMGIPTFGR
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                                                                                                                                                                                           Parekh RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptide; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy
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                                                                                                                                                                                           Rohlff C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2;
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Pred. No. 0.
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0.99;
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The invention relates to screening, diagnosis or prognosis of Vascular Dementia (VD) in a subject comprising analysing body fluid from the subject by 2-dimensional (2-D) electrophoresis to generate a 2-D array

a 2-D array of

Claim 6; Page 36; 151pp; English.

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24-NOV-2000;
28-NOV-2000;
The invention relates to screening, diagnosis or prognosis of Vascular Dementia (VD) in a subject comprising analysing body fluid from the subject by 2-dimensional (2-D) electrophoresis to generate a 2-D array of features containing at least one chosen feature whose relative abundance correlates with the presence, absence, stage or severity of VD or predicts the onset or course of VD, especially detecting in a sample of cerebrospinal fluid (CSF) from the subject one of 223 VD-associated protein isoforms (VPIs) (ABB55801-ABB56295) as fully defined in the specification. Detecting VD-associated features and VPI is useful for the screening, diagnosis or prognosis of VD, for determining the stage or severity of VD, for identifying a subject at risk of VD or for monitoring the effect of therapy administered to a subject having VD. Nucleic acids
                                                                                                                                                                                                                                                                         Screening, diagnosis or prognosis of vascular dementia (VD), use determining stage of VD and monitoring the effect of VD therapy, comprises analyzing body fluid by 2-dimensional electrophoresis features correlated with VD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    features containing at least one chosen feature whose relative abundance correlates with the presence, absence, stage or severity of VD or predicts the onset or course of VD, especially detecting in a sample of cerebrospinal fluid (CSF) from the subject one of 223 VD-associated protein isoforms (VPIs) (ABB55801-ABB56295) as fully defined in the specification. Detecting VD-associated features and VPI is useful for the screening, diagnosis or PVD-associated features and vPI is useful for severity of VD, for identifying a subject at risk of VD or for monitoring the effect of therapy administered to a subject having VD. Nucleic acids encoding a VPI or inhibiting the function of a VPI are useful for the
                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-557937/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vascular dementia-associated protein isoform (VPI) 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-FEB-2002
                                                                                                                                                                                                                                       Claim 6; Page 32; 151pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200169261-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vascular Dementia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treatment of VD and for gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                      HMAC,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; peptide;
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2000US-00724391.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene
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Pred. No. 0.99
2; Mismatches
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RESULT 9
AAU28751
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Best Local
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08-DEC-2000;
12-DEC-2000;
                                    The present invention relates to the identification of depression associated protein isoforms (DPIs), particularly the tryptic digest peptides of these proteins. Some of the DPIs (AAU28404-AAU28625) described are decreased in the cerebrospinal fluid (CSF) of BAD (bip affective disorder) subjects, whilst other DPIs (AAU28626-AAU28887) increased in BAD subjects. Also described are peptide sequences identified from DPI-45 and DPI-213 and the nucleic acid sequence the encoded by. The sequences of the invention are useful for clinical screening, diagnosis, prognosis, therapy and prophylaxis of neuropsychiatric disorders e.g. BAD (also known as bipolar mood diso BP), maniac-depressive illnesses, attention deficit disorders. The presequence represents one of the DPI tryptic digest peptides of the presequence represents one of the DPI tryptic digest peptides
                                                                                                                                                                                                                                                               disorder, which i
neuropsychiatric
                                                                                                                                                                                                                                                                                                                        WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; depression associated protein isoform; tryptic digest peptide; DPI; cerebrospinal fluid; CSF; BAD; bipolar affective disorder; neuropsychiatric disorder; bipolar mood disorder; neuroleptic; maniac-depressive illness; schizoaffective disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DPI tryptic digest peptide #348.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU28751 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      encoding a VPI or inhibiting the function of a VPI are useful for the treatment of VD and for gene therapy
                                                                                                                                                                                                                                    Disclosure; Page 38; 153pp; English
                                                                                                                                                                                                                                                                                            Novel
                                                                                                                                                                                                                                                                                                                                                  Herath HMAC,
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                                                                                                                                                                                                                                                                                                                        2001-570626/64
                                                                                                                                                                                                                                                              nucleic acid encoding der, which is used for psychiatric disorders,
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2000GB-00030050
2000US-0254830P
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                                                                                                                                                                                                                                                                                                                                                Rohlff C,
                                                                                                                                                                                                                                                            a protein associated with bipolar affective diagnosis, prophylaxis and therapy of such as bipolar affective disorder.
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Pred. No. 0.99
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                  Terrett
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1.99;
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Query Match Best Local S Matches

Similarity 70.

49.4%;

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Score 41; DB 4 Pred. No. 0.99; 2; Mismatches

4.

Length 10

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Sequence 10

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RESULT 11
AAU28539
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                                                                                                                               Matches
                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                   associated protein isoforms (DPIs), particularly the tryptic digest peptides of these proteins. Some of the DPIs (AAU28404 AAU28625) described are decreased in the cerebrospinal fluid (CSF) of BAD (bipolar affective disorder) subjects, whilst other DPIs (AAU28626-AAU28887) are increased in BAD subjects. Also described are peptide sequences identified from DPI-45 and DPI-213 and the nucleic acid sequence they are encoded by. The sequences of the invention are useful for clinical screening, diagnosis, prognesis, therapy and prophylaxis of neuropsychiatric disorders, e.g. BAD (also known as bipolar mood disorder, neuropsychiatric disorders, e.g. BAD (also known as bipolar mood disorder, neuropsychiatric disorders, e.g. BAD (also known as bipolar mood disorder).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel nucleic acid encoding a protein associated with bipolar affective disorder, which is used for diagnosis, prophylaxis and therapy of neuropsychiatric disorders, such as bipolar affective disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; depression associated protein isoform; tryptic digest peptide; DPI; cerebrospinal fluid; CSF; BAD; bipolar affective disorder; neuropsychiatric disorder; bipolar mood disorder; neuroleptic;
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08-DEC-2000; 2000GB-00030050.
12-DEC-2000; 2000US-0254830P.
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                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                        BP), maniac-depressive illnesses, attention deficit disorders, schizoaffective disorders, and unipolar affective disorders. The present sequence represents one of the DPI tryptic digest peptides of the presen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
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                                                                                            w
                                                                                                                                             Similarity
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                                                                LVMGIPTFGR 10
                                                                                            LVMGVPFYGR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                   invention relates to the identification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 38; 153pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 digest peptide #388.
                                                                                                                                                                                           AA;
                                                                                                                              Conservative
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70.0%;
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                                                                                                                                             Score 41; I
Pred. No. 0.
                                                                                                                              Mismatches
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                                                                                                                                             DB 4;
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RESULT 12
AAU25095
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AC AAU25
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                                                                                                                                                                                                                                                                                                                                                     associated protein isoforms (DPIs), particularly the tryptic digest peptides of these proteins. Some of the DPIs (AAU28404-AAU28625) described are decreased in the cerebrospinal fluid (CSF) of BAD (bipolar affective disorder) subjects, whilst other DPIs (AAU28626-AAU28887) are increased in BAD subjects. Also described are peptide sequences identified from DPI-45 and DPI-213 and the nucleic acid sequence they are encoded by. The sequences of the invention are useful for clinical screening, diagnosis, prognosis, therapy and prophylaxis of neuropsychiatric disorders e.g. BAD (also known as bipolar mood disorder, BP), maniac-depressive illnesses, attention deficit disorders. The present sequence represents one of the DPI tryptic digest peptides of the present represents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel nucleic acid encoding a protein associated with bipolar affective disorder, which is used for diagnosis, prophylaxis and therapy of neuropsychiatric disorders, such as bipolar affective disorder.
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08-DEC-2000; 2000GB-00030050.
12-DEC-2000; 2000US-0254830P.
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                                                                                                                                                                                                                                                                                                                Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
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                Schizophrenia-Associated Protein Isoform
                                                18-DEC-2001
                                                                                                           AAU25095 standard; peptide; 10
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                                                                                                                                                                                                                                                                                                                A.
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                                             (first entry)
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                                                                                                                                                                                                                                                                 49.48;
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                                                                                                                                                                                                                                                                 Score 41; I
Pred. No. 0.
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0.99;
                   (SPI) peptide #324
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RESULT 13
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Best Local
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 24-FEB-2000;
08-DEC-2000;
12-DEC-2000;
                                                                                                                                                                        Human; Bipolar Affective Disorder; BAD; Depression-Associated feature; DF; Depression-Associated protein isoform; DR]; Cerebro-spinal fluid; CSF; antidepressant; antimanic; nootropic; tranquiliser; neuroleptic; attention deficient disorder; schizoaffective disorder;
                                                                                                                                                                                                                                                                     18-DEC-2001
                                                                                                                                                                                                                                                                                                                       AAU26399 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       These protein isoforms, e.g. SPI-206, SPI-238 and SPI-240 are detectable in cerebrospinal fluid, serum or plasma and are useful markers of schizophrenia. The sequences can be used for treatment and diagnosis of schizophrenia, screening, prognosis, monitoring the results of therapy, identifying patients most likely to respond to a particular therapy and identification of new targets for drug treatment. SPI DNA is useful as a nucleic acid probe to detect the presence of nucleic acids or SPIs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New schizophrenia associated protein isoforms and encoding nucleic acid molecules, useful for treatment, diagnosis and prognosis of schizophrenia and screening for potential drugs for treatment and new drug targets.
                                                    23-FEB-2001;
                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schizophrenia-associated protein isoform; SPI; SPI-206; SPI-238; SPI-240; neuroleptic; gene therapy; cerebrospinal fluid; serum; plasma.
                                                                                                         WO200163294-A2
                                                                                                                                                             unipolar affective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10
                                                                                                                                                                                                                                                                                                                                                                                                                   LVMGVPFYGR 12
                                                                                                                                                                                                                                                                                                                                                                                          LVMGIPTFGR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 35; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
2000GB-00004412.
2000GB-00030050.
2000US-0254830P.
                                                    2001WO-GB000791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      represents a schizophrenia-associated protein isoform (SPI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000GB-00004415.
2000US-00750395.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001WO-GB000792
                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                             disorder
                                                                                                                                                                                                                                                                                                                                                                                          10
                                                                                                                                                                                                                                                                                                                                                                                                                                                          49.4%;
                                                                                                                                                                                                                                         Protein isoform DPI-184 #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rohlff C,
                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 41; DB Pred. No. 0.99
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            . 99 ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 10
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RESULT 14
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XX AAU26
XX Depre
XX Depre
XX Human
KW DF; F
KW CSF;
KW CSF;
KW unipc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   used to screen, diagnose or prognose of BAD or unipolar depression, determine the stage or severity of BAD or unipolar depression, identify a subject at risk of developing BAD or unipolar depression, or monitor the effect of therapy in a subject. They are also used to screen for or identify agents that interact with a DPI. These agents, antibodies agents the DPIs, and nucleic acids encoding the DPIs are used to treat or prevent BAD or unipolar depression. Diseases that can be treated are attention deficient disorder, a schizoaffective disorder, a bipolar or a unipolar affective disorder. The DPIs are used in proteomics. The proteomic approach of using DPIs for screening, diagnosis or prognosis of EAD or unipolar depression overcomes the problems of using gene expression analysis, such as not being able to obtain central nervous system (CNS) tissue from a living patient under normal circumstances. The present sequence is a DIP increased in the CSF (cerebro-spinal fluid) of
                                                                                                                     24-FEB-2000; 2000GB-00004412
08-DEC-2000; 2000GB-00030050
12-DEC-2000; 2000US-0254830P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; Bipolar Affective Disorder; BAD; Depression-Associated feature; DF; Depression-Associated protein isoform; DPI; Cerebro-spinal fluid; CSF, antidepressant; antimanic; nootropic; tranquiliser; neuroleptic; attention deficient disorder; schizoaffective disorder;
                                                            (OXFO-) OXFORD GLYCOSCIENCES UK LTD
                                                                                                                                                                                                                                                                                                              30-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                       WO200163294-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             unipolar affective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAU26439 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Preparation for diagnosing or treating bipolar affected disorder (BAD) unipolar depression, or for screening for modulators, comprises a BAD-
                                                                                                                                                                                                                                                  23-FEB-2001; 2001WO-GB000791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Depression-Associated Protein isoform DPI-234.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                subjects having BAD
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                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens
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ed protein isoform.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10
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70.0%;
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Pred. No. 0.
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Mismatches
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Herath HMAC,

Parekh

RB,

Rohlff C

WPI;

2001-582081/65

Preparation for diagnosing or treating bipolar affected disorder (BAD) unipolar depression, or for screening for modulators, comprises a BAD-

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RESULT 15
AAU26186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a preparation comprising an isolated Bipolar Affected Disorder (BAD)-Associated Protein Isoform (DFIs). The DPI's are used to screen, diagnose or prognose of BAD or unipolar depression, identify a determine the stage or severity of BAD or unipolar depression, identify a subject at risk of developing BAD or unipolar depression, or monitor the effect of therapy in a subject. They are also used to screen for or conditivity agents that interact with a DPI. These agents, antibodies against the DPIs, and nucleic acids encoding the DPIs are used to treat or prevent BAD or unipolar depression. Diseases that can be treated are attention deficient disorder. The DPIs are used in proteomics. The DPIs are used in proteomics of continuous affective disorder. The DPIs are used in proteomics of DPIs are used in proteomics. The DPIs are used in proteomics. The proteomic approach of using DPIs for screening, diagnosis or prognosis of CR Spites (CNS) tissue from a living patient under normal circumstances. The present sequence is a DIP increased in the CSF (cerebro-spinal fluid) of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unipolar depression, or for associated protein isoform.
Preparation for diagnosing
                                 WPI; 2001-582081/65
                                                                                                                             24-FEB-2000; 2000GB-00004412.
08-DEC-2000; 2000GB-00030050.
12-DEC-2000; 2000US-0254830P.
                                                                                                                                                                                                                                 30-AUG-2001.
                                                                                                                                                                                                                                                                 WO200163294-A2
                                                                                                                                                                                                                                                                                                                                unipolar affective disorder.
                                                                                                                                                                                                                                                                                                                                              DF; Depression-Associated protein isoform; DPI; Cerebro CSF; antidepressant; antimanic; nootropic; tranquiliser attention deficient disorder; schizoaffective disorder;
                                                                                                                                                                                                                                                                                                                                                             Human; Bipolar Affective Disorder; BAD; Depression-Associated feature DF; Depression-Associated protein isoform; DPI; Cerebro-spinal fluid; CSF; antidepressant; antimanic; nootropic; tranquiliser; neuroleptic;
                                                                                                                                                                                                                                                                                                                                                                                                                             Depression-Associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU26186 standard; peptide; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 8; Page 38; 163pp; English.
                                                                                                                                                                                                 23-FEB-2001; 2001WO-GB000791.
                                                                                                  (OXFO-)
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                                                                                                  OXFORD GLYCOSCIENCES UK LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                  Parekh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             49.4%;
                                                                RВ,
                                                                                                                                                                                                                                                                                                                                                                                                                             Protein isoform DPI-49
                                                                Rohlff
or treating bipolar affected disorder (BAD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 41; DB 4
Pred. No. 0.99;
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unipolar depression, or for associated protein isoform.
                                                                                                                                     The invention relates to a preparation comprising an isolated Bipolar Affected Disorder (BAD)-Associated Protein Isoform (DPIs). The DPI's
                                                                                                                                                                                       or for screening
                                                                                                                                                              English
                                                                                                                                                                                        for modulators, comprises a BAD-
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used to screen, diagnose or prognose of BAD or unipolar depression, identify a content of therapy in a subject. They are also used to screen developing BAD or unipolar depression, identify a content of the party of the prognose of BAD or unipolar depression, identify a content of therapy in a subject. They are also used to screen for or content of the DPIs, and nucleic acids encoding the DPIs are used to treat corprevent BAD or unipolar depression. They are also used to screen for or content of the DPIs, and nucleic acids encoding the DPIs are used to treat corprevent BAD or unipolar depression. Diseases that can be treated are content of disorder, a schizoffective disorder, a bipolar or a compolar affective disorder. The DPIs are used in proteomics. The composition analysis, such as not being able to obtain central nervous content sequence is a DIP decreased in the CSF (cerebro-spinal fluid) of subjects having BAD The of of. þ

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Matches
                                           Query Match
Best Local
                                          Local
               3 LVMGVPFYGR 12
                                 Similarity 7; Conserv
||||:| :||
LVMGIPTFGR 10
                                  Conservative
                                          49.4%;
70.0%;
                                           Score 41; I
Pred. No. 0.
                                  Mismatches
                                                   DB 4;
                                  1;
                                                   Length 10;
                                  Indels
                                  0
                                  Gaps
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Sequence 10 AA;

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Homo sapiens.
24-FEB-2000; 2000GB-00004415.
28-DEC-2000; 2000US-00750395.
                                                                          23-FEB-2001; 2001WO-GB000783
                                                                                                                             30-AUG-2001
                                                                                                                                                                           WO200163293-A2
                                                                                                                                                                                                                                                                          neurological disorder; neuropathy.
                                                                                                                                                                                                                                                                                                 Schizophrenia; neuroleptic; diagnostic; neuropsychiatric disorder;
                                                                                                                                                                                                                                                                                                                                                     Schizophrenia-associated isoform peptide #324.
                                                                                                                                                                                                                                                                                                                                                                                                     24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU15439 standard;
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide; 10
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The invention relates to methods and compositions for screening, diagnosis and prognosis of Schizophrenia. The method involves de the presence of Schizophrenia (SCH) Associated Features (SFs) are

detecting

Page

35; 160pp; English.

Diagnosing and monitoring Schizophrenia by detecting the presence of Schizophrenia Associated Protein Isoforms in samples of cerebrospinal fluid.

WPI; 2001-502868/55.

(OXFO-) OXFORD GLYCOSCIENCES UK LTD

Parekh

RВ,

Rohlff C;

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RESULT 17
ABB52306
ID ABB52
XX ABB52
XX ABB52
XX ABB52
XX ABB52
XX Human
XX Human
XX Homo
XX Expre
XX Alzhe
XX Homo
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC Associated Protein Isoforms (SPIs) in samples, e.g. by electrophoresis, communoassay or hybridisation assay, for diagnosing and monitoring SCH, communoassay or hybridisation assay, for diagnosing and monitoring SCH, communoassay or hybridisation assay, for diagnosing and monitoring SCH, communoassay of the effectiveness of treatments and for identifying potential communoassay and the relative abundance of at least 1 chosen feature or diagnosis of CC SCH and the presence or absence of SCH; and (2) monitoring the effect of CC at least 1 chosen feature which correlates with the severity of SCH. The CC expression and activity of the SFS, SPIs and related molecules (e.g. condary messengers) are studied to diagnose SCH, monitor the progress CC identify and produce potential therapeutic agents for the treatment of cultural rand as targets to CC identify and produce potential therapeutic agents for the treatment of neuropsychiatric disorders, where CC manifestations of anatomical and biochemical changes have been identified CC in many cases. Consequently the identification and characterisation of CC cellular and/or molecular causative defects and neuropathies are CC necessary for improved treatment of neuropsychiatric disorders. ANU15114- and/or molecular causative defects and neuropathies are CC necessary for improved treatment of neuropsychiatric disorders. ANU15114- and/or molecular causative defects and neuropathies are CC necessary for improved treatment of neuropsychiatric disorders.
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Best Local
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                                                                              Screening for Alzheimer's disease in a mammal, by making two-dimensional array of a feature whose relative abundance correlates with disease, and comparing with abundance of the feature in samples of healthy persons.
                                                                                                                                                                                                                                                                                                          Durham
Potter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-APR-2000;
28-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; neuroprotective; nootropic; gene therapy; vaccine; Alzheimer's disease; Alzheimer's Disease-Associated Feature; Alzheimer's Disease-Associated Protein Isoform; API; tryptic Expression Reference Protein Isoform; ERPI; proteolysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human API-41 tryptic
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                                                                                                                                                                                                           WPI; 2001-639384/73
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                                                                                                                                                                                                                                                                                                                                                                                           OXFORD GLYCOSCIENCES UK LTD PFIZER INC.
                                                                                                                                                                                                                                                                       RR,
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Rohlff C,
R, White F,
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2000US-0253647P
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                                                                                                                                                                                                                                                                    T. DL, Herath HMAC
C, Silber BM, St
F, Williams SA;
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Pred. No. 0.99;
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                HMAC,
                                                                                                                                                                                                                                                                                                MAC, Kimmel
Stiger TR,
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Sunderland D
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Example; Page 33; 162pp; English

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Query Match

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Best Local
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                                     The present invention relates to methods for screening or diagnosing Alzheimer's disease (AD) to determine the stage or severity of AD in subject, to identify subject at risk of developing AD, or to monitor effect of therapy administered. The methods comprise analysing a test sample of body fluid by 2-dimensional electrophoresis to generate a 2 dimensional array of AD-associated features (AFS). The method alternatively comprises quantitatively detecting in a sample of body fluid from the subject, one or more AD-associated protein isoforms (A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to methods for the screening, diagnosis and prognosis of Alzheimer's disease. The methods involve the detection of Alzheimer's Disease-Associated Features (AFS) and Alzheimer's Disease-Associated Protein Isoforms (APIS) in cerebrospinal fluid, serum or plasma. The abundance of the AFS and APIS is then normalised to an Expression Reference Protein Isoform (ERPI) in order to determine whether a patient is suffering from, or has a predisposition to, Alzheimer's Disease. The relative abundance of the AFS and APIS correlates with the severity of Alzheimer's Disease. The present sequence is a peptide produced from an API by proteolysis
                                                                                                                                                                                                   Screening or diagnosing of Alzheimer's disease (AD) determine or severity of AD in a subject, comprises analyzing a test sam fluid from the subject by 2-dimensional electrophoresis.
                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                 WPI; 2003-371957/35
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                           ABR58710-ABR59184)
                                                                                                                                                                                                                                                                                              Sunderland PT,
                                                                                                                                                                                                                                                                                                              Potter DM,
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                                                                                                                                                                        2; Page 51; 179pp; English.
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7; Conserva
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OXFORD GLYCOSCIENCES UK LTD.
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LVMGIPTFGR
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Rohlff C, Silber BM, Snyd
PT, Townsend RR, White WF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disease-associated protein isoform
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Pred. No. 0.99
2; Mismatches
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                                                                                                                                                                                                                                                                                             Snyder PJ, Some WF, Williams
                                                                                                                                                                                                                                                                                                                    Kimmel LH,
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ams SA;
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of body
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                                                                                                                                               The invention relates to a novel method for screening or diagnosis of Alzheimer's disease (AD) in a subject, determining the stage or severity of AD, identifying a subject at risk of developing AD or monitoring the effect of therapy administered to a subject having AD, by analysing body fluid to generate a two-dimensional array of Alzheimer's disease-associated features (AFs) such as AF-200, AF-201, AF-202, AF-203, AF-204, AF-205, etc., and comparing the abundance of AFs with a control. The method of the invention has nootropic and neuroprotective applications and may be useful for screening or diagnosis of Alzheimer's disease (AD) in a subject, determining the stage or severity of AD in a subject, determining the stage or severity of AD in a subject, identifying a subject at risk of developing AD or monitoring the effect of therapy administered to a subject having AD. The body fluid is cerebrospinal fluid (CSF). The current sequence is that of a human Alzheimer's disease-associated protein isoform (API) tryptic digest
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Matches
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Best Local :
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                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                           disease-Associated Features.
                                                                                                                                                                                                                                                                                                                                                                                                     Screening or diagnosis of Alzheimer's disease (AD) in subject, determining stage or severity of AD, identifying subject at risk of developing AD, or monitoring effect of therapy, by detecting Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-318939/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Durham LK,
Potter DM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CSF; Alzheimer's disease-associated protein isoform; API; tryptic digest;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alzheimer's disease; nootropic; neuroprotective; cerebrospinal fluid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Alzheimer's disease-API tryptic digest peptide -
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                                                     Similarity 7; Conserv
                                                                                                                                        of the
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OXFORD GLYCOSCIENCES UK LTD
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 LVMGIPTFGR 10
                           LVMGVPFYGR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Friedman DL, Herath HMAC, Kimmel LH, Rohlff C, Silber BM, Snyder PJ, Soa
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                                                        Conservative
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                                                                                                                                        invention.
                                                                                                                                                                                                                                                                                                                                                               277; 208pp; English.
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2; Mismatches
                                                                    Score 41;
Pred. No.
                                                        Mismatches
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Query Match Best Local S Matches 7

Similarity 7; Conserv

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49.4%;

Score 41; DB 8 Pred. No. 0.99;

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Length 10;

Mismatches

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ID AD0788
XX AD078
AC AD078

                                                                              The invention describes an isolated nucleic acid molecule (I) that CC hybridises to two short nucleic acid sequences and the 1515 amino acid compared to the specification. Also described are: a CC preparation comprising an isolated peptide coded for by the nucleic acid molecule above, or comprising an isolated human protein comprising one or CC more of the following sequences: Glu-Leu-App-Val-Leu-Gly-Arg; methods for CC diagnosing Schizophrenia; antibodies capable of immunospecific binding to a Schizophrenia; and methods of screening for agents that modulate a CC chizophrenia; and methods of screening for agents that modulate a CC characteristic (e.g., expression or binding activity) of an SPI and CC canalogue, or an SPI-related polypeptide. The nucleic acid molecule and concoded proteins, as well as the methods and compositions are useful for screening, diagnosing, and prognosing Schizophrenia, for monitoring the effectiveness of Schizophrenia treatment, for identifying patients most CC developing drug. They are also useful for screening modulators of Schizophrenia. This is the amino acid sequence a schizophrenia-associated protein isoform increased in the cerebrospinal fluid of schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated nucleic acid molecule, useful for dia
for monitoring the effectiveness of Schizophrenia
screening agents for treating Schizophrenia.
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28-DEC-2000;
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Schizophrenia-Associated Protein Isoform; SPI; schizophrenia screening
schizophrenia diagnosing; schizophrenia prognosing;
Schizophrenia treatment; drug development; cerebrospinal fluid; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-AUG-2004
   Sequence
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TYSON K L.
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   ΑA,
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2000US-00750395.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 spi; schizophrenia screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ú
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosing Schizophrenia, nia treatment or for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Terrett
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                 Matches
                         Query Match
Best Local
                                                 Sequence 10 AA;
                        Local
3 LVMGVPFYGR 12
                Similarity 7; Conserv
                Conservative
                       49.4%;
                        Score 41; I
Pred. No. 0.
                Mismatches
                        DB 8;
                μ.
                               Length 10;
                Indels
                0
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RESULT 23
ADV22957
ID ADV7
XX
AC AT
XX
DT
XX
XX
XX

ADV22957 standard;

peptide; 18

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10-MAR-2005 ADV22957;

(first entry)

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IPFYGKAIPLE

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RESULT 21
ADR19350
ID ADR1
                          The invention relates to a novel method for assessing, screening, containing or prognosing chronic tissue damage in a subject identifying the effect of therapy administered to a subject having chronic tissue damage. Or monitoring the content of therapy administered to a subject having chronic tissue damage. Content tissue damage, or tissue damage, or monitoring the content of the subject a tissue repair-associated protein isoform (TRPI) given the specification. The invention further comprises an isolated or recombinant TRPI selected from the proteins listed in the specification; content fixed an agent capable of modulating the correction or activity of the TRPI or a TRPI related polypeptide; a composition comprising the TRPI cited above or an active agent that modulates the expression and/or activity of the TRPI or a TRPI cated polypeptide, and a pharmaceutical carrier, vehicle or diluent; conducted polypeptide, and a pharmaceutical carrier, vehicle or diluent; conducted polypeptide, and a pharmaceutical carrier, vehicle or diluent; conducted polypeptide, and a pharmaceutical carrier, vehicle or diluent; conducted polypeptide, and a pharmaceutical carrier, vehicle or diluent; conducted polypeptide, and a pharmaceutical carrier, vehicle or diluent; conducted polypeptide, prognosing and treating chronic tissue damage, conducted the processing and treating chronic tissue damage, conducted the activity of TRPI or TRPI related polypeptide. The conductation are produced by proteolysis using tryptic digest peptide. The conductation derived human tryptic digest peptide. The conductation isoform of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Butt R, AL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diagnosing, or monitoring the effect of therapy of, chronic tissue (e.g. chronic dermal ulcer) comprises detecting and/or quantifying Repair-Associated protein isoform in a sample (e.g. wound exudate)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 7; SEQ ID NO 12; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-604404/58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PFIZ )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-FEB-2003; 2003US-0444600P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-FEB-2004; 2004WO-US002881.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue damage; chronic; tissue repair-associated protein isoform; TRPI; antiulcer; vulnerary; chronic dermal ulcer; human; tryptic digest;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TRPI derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            proteolysis;
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OXFORD GLYCOSCIENCES UK LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rumpel K,
, Patel TP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tryptic digest
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Williams-Jones BI,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rohlff C;
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Tissue
from
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RESULT 22
ADL26337
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                           Matches
                                        Query Match
Best Local
                                                                                                     The invention relates to a novel method for isolating Hepatitis C Virus (HCV) peptides (HPs). The method of the invention has virucide activity, and may be useful in producing a vaccine. The method is useful for isolating Hepatitis C Virus peptides (HPs) which have a binding capacity to a MHC/HLA molecule or a complex comprising the HCV-peptide and the MHC/HLA molecule for preparing a vaccine against HCV infection. The Teclls, a T cell clone or a Tepperaring a composition for identifying heteroclitic epitopes or for preparing a composition for treating HCV infection. The present sequence represents a synthetic peptide derived from a conserved region of HCV.
                                                                               Sequence 15
                                                                                                                                                                                                                                                                                   molecule by separating bind to the molecule.
                                                                                                                                                                                                                                                                                             Isolating Hepatitis C Virus peptides (HPs) which have a binding capacity to a MHC/HLA molecule or a complex comprising the HCV-peptide and the molecule by separating the complex from the HCV-peptides which do not
                                                                                                                                                                                                                                                                                                                                                                                        Buschle M,
                                                                                                                                                                                                                                                                                                                                                                                                                                             13-SEP-2002; 2002AT-00001376
27-FEB-2003; 2003WO-EP002005
11-JUL-2003; 2003EP-00450171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADL26337 standard; peptide;
                                                                                                                                                                                                                                                        Example 1; Page 31;
                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-269899/25.
                                                                                                                                                                                                                                                                                                                                                                             Zauner W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-AUG-2003; 2003WO-EP009482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HCV; hepatitis C virus; virucide; vaccine; MHC; HLA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         histocompatibility
 7 VPFYGRAXSIE 17
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6; Conserv
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LVMGIPTFGR
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Zinke S,
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                           Conservative
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                                                                                                                                                                                                                                                       73pp;
                                      48.2%;
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Kirlappos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         complex; human leukocyte
                                                                                                                                                                                                                                                      English.
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                                      Score 40; DB
Pred. No. 2.4;
                                                                                                                                                                                                                                                                                                                                                                            Mattner F, H;
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                           Mismatches
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                           <u>۲</u>
                                                    Length 15;
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                           Gaps
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Vaccine; virucide;

antigen; autoimmune disease; infection;

H77 immunogenic peptide

#198

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CC peptides above and thus presents the peptides, a population of such CC antigen-presenting cells, a process for producing antigen-presenting cells for modulating an immune response to a polypeptide of interest, a CC method for producing antigen-specific lymphocytes, a composition CC comprising at least one set of the peptides (and a carrier and/or CC diluent), a method for modulating an immune response to a polypeptide of interest comprising administering to a patient in need at least one set of the peptides, a method for treatment and/or prophylaxis of a disease or condition associated with the presence of a polypeptide of interest and composition of matter for modulating an immune response in a cc subject to a target antigen. The polypeptide of interest is also a CC disease - or condition-associated polypeptide that is a polypeptide of graduced by a pathogenic organism or a cancer, and produced by a pathogenic organism or a cancer, and produced by a pathogenic organism selected from yeast, viruses, bacteria, helminths, CC polypeptide is produced by a cancer selected from melanoma, lung cancer, cancer, stomach cancer, bladder cancer, kidney cancer, post transplant CC lymphoproliferative disease (PTLD) or Hodgkin's Lymphoma. The uncultured antigen-presenting cells or their precursors are useful in the condition is associated with the presence or condition is associated with the presence or condition associated or their precursors are useful in the condition is associated with the presence or condition is associated with the presence or
                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                            in a subject, which disease or condition is associated with the presence or aberrant expression of a target antigen, where the antigen-presenting cells or their precursors have not been subjected to activating conditions but have been contacted with an antigen that corresponds to the target antigen to express a processed or modified form of the antigen for presentation to the subject's immune system. The present sequence is one of a set of overlapping immunogenic peptides derived from a Hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to the use of at least one set of peptides in the preparation of a medicament for modulating an immune response, where individual peptides of a respective set comprise different portions of an amino acid sequence corresponding to a single polypeptide of interest and display partial sequence identity or similarity to at least one other peptide of the same set of peptides (i.e. they are overlapping). Also included are an antigen-presenting cell which has been contacted with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Use
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                                                                                                                                                                                                                  Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 1377; 645pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               viral, bacterial, protozoal and mycoplasma
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25-MAR-2004;
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Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                     protein
                                                                                                                                                                                                                  B
           Conservative
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2004AU-00901589.
                                                 48.2%;
Score 40; DB Pred. No. 3; Mismatches
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RESULT 24
ADV22958
ID ZDV2
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XX
ADV2
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ADV2
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ADV2
XX
Vacc
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                                                                   cc peptides above and thus presents the peptides, a population of such cantigen-presenting cells, a process for producing antigen-presenting cells for modulating an immune response to a polypeptide of interest, a cc method for producing antigen-specific lymphocytes, a composition comprising at least one set of the peptides (and a carrier and/or cd diluent), a method for modulating an immune response to a polypeptide of interest comprising administering an immune response to a polypeptide of interest control of the peptides, a method for treatment and/or prophylaxis of a disease cor condition associated with the presence of a polypeptide of interest can a composition of matter for modulating an immune response in a cc subject to a target antigen. The polypeptide of interest is also a disease or condition-associated polypeptide of interest is also a concease of a pathogenic organism or a cancer, and produced by a pathogenic organism or a cancer, and produced by a polypeptide is produced by a cancer selected from yeast, viruses, bacteria, helminths, polypeptide is produced by a cancer selected from melanoma, lung cancer, corvictal cancer, prostate cancer, post transplant cancer, stomach cancer, bladder cancer, kidney cancer, post transplant cancer, stomach cancer, bladder cancer, kidney cancer, post transplant cancer, their response to a polypeptide of the cancer cancer cancer, pacereatic cancer, bladder cancer, kidney cancer, post transplant cancer, carvictal cancer, prostate cancer, post transplant cancer, cancer canc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preparation of a medicament for modulating an immune response, where individual peptides of a respective set comprise different portions of an amino acid sequence corresponding to a simple polypeptide of interest and display partial sequence identity or similarity to at least one other peptide of the same set of peptides (i.e. they are overlapping). Also included are an antigen-presenting cell which has been contacted with the
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breast tumor; uterine cervix tumor; prostatic cancer; colon tumor;
pancreas tumor; stomach tumor; bladder tumor; kidney tumor;
lymphoproliferative disease (PTLD) or Hodgkin's Lymphoma. antigen-presenting cells or their precursors are useful i preparation of a medicament for the treatment of a diseas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of at least one set of peptides in the preparation of a medicament for modulating an immune response, and for treating cancer or yeast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUN-2003;
25-MAR-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 1378; 645pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 viral, bacterial, protozoal and mycoplasma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HCV H77 immunogenic peptide #199.
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2004AU-00901589.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         relates to the use of at least one set of peptides in the
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    recursors are useful in treatment of a disease
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RESULT 25
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            The present invention relates to methods for screening or diagnosing Alzheimer's disease (AD) to determine the stage or severity of AD in a subject, to identify subject at risk of developing AD, or to monitor the effect of therapy administered. The methods comprise analysing a test gample of body fluid by 2-dimensional electrophoresis to generate a 2-dimensional array of AD-associated features (AFS). The method alternatively comprises quantitatively detecting in a sample of body fluid from the subject, one or more AD-associated protein isoforms (APIs, and a sample of body comprises controlled from the subject, one or more AD-associated protein isoforms (APIs, and a sample of body comprises).
                                                                                                                                                                                                                                                                                                                                                  Screening or diagnosing of Alzheimer's disease (AD) determine the or severity of AD in a subject, comprises analyzing a test sample fluid from the subject by 2-dimensional electrophoresis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in a subject, which disease or condition is associated with the presence or aberrant expression of a target antigen, where the antigen-presenting cells or their precursors have not been subjected to activating conditions but have been contacted with an antigen that corresponds to the target antigen to express a processed or modified form of the antigen for presentation to the subject's immune system. The present sequence is one of a set of overlapping immunogenic peptides derived from a Hépatitis
                                                                                                                                                                                                                                                                                             Claim 2; Page 47; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-371957/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sunderland
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Potter DM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alzheimer's Disease-associated protein isoform, API-406, SEQ ID 278
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PFIZ ) PFIZER PROD INC. (OXFO-) OXFORD GLYCOSCIENCES UK LTD.
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ABR58710-ABR59184)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Friedman DL, Herath HMAC, Kimmel LH, Parekl Rohlff C, Silber BM, Snyder PJ, Soares HD, PT, Townsend RR, White WF, Williams SA;
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erem HD, Stiger TR;
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2: /cgn2_6/ptodata/1/iaa/f_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

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                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: DERIVED FROM ; OTHER INFORMATION: SEQUENCE OF HUMAN CARTILAGE (HC) -39 PROTEIN US-09-171-705-39
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US-09-171-705-39
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APPLICANT: BOOTS, ANNA M.H.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR USE IN ANTIGEN SPECIFIC TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY
FILE REFERENCE: 0/96198 US
CURRENT APPLICATION UNWEER: US/09/171,705
CURRENT FILING DATE: 1999-02-09
NUMBER OF SEQ ID NOS: 78
NUMBER OF SEQ ID NOS: 78
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NUMBER OF SEQ ID NOS: 49
SOFTWARE: PACENTIN Ver. 2.0
SEQ ID NO 10
LENGTH: 17
TYPE: PRT
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Patent No. 6184204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 39
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                                                                                                               Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Dermatophagoides farinae FEATURE: OTHER INFORMATION: At location 14,
                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
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                                                KLVMGVPFYGRA 13
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                                                                                                                                                    56.6%;
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                                                                                                                                                    Score 47; DB 2;
Pred. No. 0.037;
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1.7e-08;
                                                                                                                   1; Indels
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US-09-171-705-40
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                                                                                     US-09-657-757-40
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                                                Sequence 40, Application US/09657757
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Best Local
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APPLICANT: VERHEIJDEN, GILBERTUS F.M.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR USI
TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY
FILE REFERENCE: 0/96198 US
CURRENT APPLICATION NUMBER: US/09/657,757
CURRENT FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: US/09/171,705
PRIOR APPLICATION NUMBER: US/09/171,705
PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 78
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 39
                                                                                                                                                                                                                                                                           APPLICANT: BOOTS, ANNA M.H.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR
TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY
FILE REFERENCE: 0/96198 US
CURRENT APPLICATION NUMBER: US/09/171,705
CURRENT FILLING DATE: 1999-02-09
NUMBER OF SEQ ID NOS: 78
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 40
LENGTH: 16
TYPE: PRT
                                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: DERIVED FROM ; OTHER INFORMATION: SEQUENCE OF HUMAN CARTILLAGE (HC) -39 PROTEIN US-09-171-705-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: DERIVED FROM
OTHER INFORMATION: SEQUENCE OF HUMAN CARTILLAGE (HC) -39 PROTEIN
US-09-657-757-39
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GENERAL INFORMATION:
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Patent No. 6184204
GENERAL INFORMATION:
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
                                        5 MGVPFYGRAXSI 16
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                                                                                   Similarity 41. 5; Conservative
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MGIPTFGRSFTL
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                                                                                                      43.4%;
41.7%;
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66.7%;
  12
                                                                                   Score 36; DB 2
Pred. No. 4.2;
5; Mismatches
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US-08-635-886C-146
; Sequence 146, Application US/08635886C
; Patent No. 6555114
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                                                       RESULT 7
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US-09-311-784A-273
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Expression Vectors for Stimulating an TITLE OF INVENTION: Immune Response and Methods of Using FILE REFERENCE: 39963-2002.01
CURRENT APPLICATION NUMBER: US/09/311,784A
CURRENT FILING DATE: 1999-05-13
PRIOR FILING DATE: 1999-05-15
PRIOR FILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 463
SOFTWARE: PastSEQ for Windows Version 3.0
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CURRENT FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: US/09/171,705
PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 78
SOFTWARE: Patentin Ver. 2.0
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APPLICANT: VERHEIJDEN, GILBERTUS F.M.
TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR USE IN ANTIGEN SPECIFIC TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY
FILE REFERENCE: 0/96198 US
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                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: DERIVED FROM
OTHER INFORMATION: SEQUENCE OF HUMAN CARTILAGE (HC) -39 PROTEIN
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nes 5; Conserv
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1 MGIPTFGRSFTL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hermanson, Gary G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Livingston,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ishioka, Glenn Y.
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71.4%;
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Pred. No. 4.2;
5; Mismatches
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Pred. No. 4.6e+05;
2; Mismatches 0;
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Best Local Similarity
Marches 5; Conserve
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                                                                                                                 RESULT 9
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; ORGANISM: hepatitis C virus
US-08-974-690C-146
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US-08-635-886C-146
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LENGTH: 8
TYPE: PRT
                                Sequence 146, Application US/08974685
Patent No. 6689368
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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Best Local Similarity 71.4%;
Matches 5; Conservative
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TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS
TITLE OF INVENTION: VIRUS
FILE REFERENCE: 2551-94
CURRENT APPLICATION NUMBER: US/08/974,690C
CURRENT FILING DATE: 1997-11-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 1997-11-19
PRIOR APPLICATION NUMBER: PCT/EP94/03555
PRIOR FILING DATE: 1994-10-28
PRIOR APPLICATION NUMBER: EP 93402718.6
PRIOR APPLICATION 1993-11-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: LEROUX-ROELS, Geert
APPLICANT: DELEYS, Robert
APPLICANT: MAERTENS, Geert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/08/635,886C CURRENT FILING DATE: 1996-04-25 PRIOR APPLICATION NUMBER: PCT/EP94/03555 PRIOR FILING DATE: 1994-10-28 PRIOR APPLICATION NUMBER: EP 3402718.6 PRIOR FILING DATE: 1993-11-04 NUMBER OF SEQ ID NOS: 286
                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: LEROUX-ROELS, APPLICANT: DELEYS, Rober APPLICANT: MAERTENS, Gee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: MAERTENS, GEET TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C TITLE OF INVENTION: VIRUS FILE REFERENCE: 2752-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
                                                                                                                                                                                                             7 VPFYGRA 13
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                                                                                                                                                                                                                                                       Conservative
  DELEYS,
                    LEROUIX-ROELS,
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                                                                                                                                                                                                                                                                        42.28;
ROBERT
                                                                                                                                                                                                                                                                        Score 35; I
Pred. No. 4
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Pred. No. 4.6e+05;
                    GEERT
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                                                                                                                                                                                                                                                                                          Length 8;
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TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES

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US-09-709-947-1
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                                                                                                                      PRIOR APPLICATION NUMBER: US 60/172,170
PRIOR FILING DATE: 1999-12-17
PRIOR PPLICATION NUMBER: US 60/178,860
PRIOR FILING DATE: 2000-01-27
PRIOR PLICATION NUMBER: US 60/201,721
PRIOR PILING DATE: 2000-05-03
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 5; Conserva
                                                                                                                                                                                                                                                                                  FILE REFERENCE: MTP-024
FILE REFERENCE: MTP-024
CURRENT APPLICATION NUMBER: US/09/709,947
CURRENT FILING DATE: 2000-11-10
PRIOR APPLICATION NUMBER: US 60/165,173
PRIOR FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09709947 Patent No. 6936424
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                                                                                                      SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Watkins, TITLE OF INVENTION: TITLE OF INVENTION:
FEATURE: OTHER INFORMATION: Description of Artificial Sequence:Tryptic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (703) 816-40
TELEFAX: (703) 816-4100
INFORMATION FOR SEQ ID NO: 146:
                                     LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/974,685
FILING DATE: 19-No. 6689368-1997
CLASSIFICATION: 4Uhknown>
ATTORNEY/AGENT INFORMATION:
NAME: BYRNE, THOMAS E.
REGISTRATION NUMBER: 32,205
REFERENCE/DOCKET NUMBER: 1487-11
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 146:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PETENTIN Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: peptid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A. ZIP: 22201-4714
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71.4%;
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Pred. No. 4.6e+05;
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Query Match
Best Local Similarity
Thes 5; Conserve
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; OTHER INFORMATION: ; OTHER INFORMATION: US-08-851-843A-171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                        TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/724
FILING DATE: 01-OCT-1996
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 0153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Andrews, William H.
TITLE OF INVENTION: No. 6093809el Telomerase
NUMBER OF SEQUENCES: 225
CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
US 08/724,643
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APPLICATION NUMBER: U
FILING DATE: 18-APR-:
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                                                                                        FEATURE:
                                                                                                     MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/0 FILING DATE: 06-MAY-1997
                                                                                                                                      TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                     LOCATION:
                                                                     NAME/KEY: Peptide
                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                           TELEPHONE:
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Nakamura, Toru
Chapman, Karen B.
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Harley, Calvin
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                                                                                                                          linear
                                                                                                                                                                                                                                             (415) 576-0200
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                   /note= "motif D peptide from human
telomerase core protein 1 (TCP1)"
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DB 2;

Length 15;

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Best Local
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PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: US 08/911,312

APPLICATION NUMBER: US 08/911,312

APPLICATION NUMBER: US 08/911,312
                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17618
PTI.ING DATE: 01-OCT-1997
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TITLE OF I
                                 ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Te
                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/844,419 FILING DATE: 18-APR-1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US OFFILING DATE: 01-OCT-1996 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy
                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                                              PRIOR APPLICATION DATA:
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                                                                APPLICATION NUMBER: FILING DATE: 01-007
                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 14-AU
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                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/846,017 FILING DATE: 25-APR-1997
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Nakamura, Karen B.
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Two Embarcadero Center, Eighth Floor
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VENTION: Human Telomerase Catalytic Subunit
EQUENCES: 727
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                                                                                                                                                                                                                                                                                                                                                           06-MAY-1997
                                                                UMBER: WO PCT/US97/17885
01-OCT-1997
                                                                                                                                                                                                           UMBER: US 08/912,951
14-AUG-1997
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77.8%;
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36,429
ER: 015389-002610US
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Best Local S
Matches 7
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INFORMATION FOR SEQ ID NO:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/854,050
FILING DATE: 09-MAY-1997
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                               APPLICATION NUMBER: US 08/846,017 FILING DATE: 25-APR-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: telomerase core protein 1 (TCP1)"
                                                                                                                   CLASSIFICATION:
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California
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INVENTION: No. 6261836el Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                    3: Townsend and Townsend and Crew LLP
Two Embarcadero Center, 8th Floor
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Nakamura, Toru
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                      United States of America
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Harley, Calvin
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77.8%;
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Pred. No. 9.
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INFORMATION:

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LOCATION: 1.15
OTHER INFORMATION: /note= "motif D peptide from human OTHER INFORMATION: telomerase core protein 1 (TCP1)" US-08-854-050-171
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US-09-430-323-171
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
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TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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Local Similarity 77.8%;
les 7; Conservarion
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LOCATION:
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REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
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                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/430,323
FILING DATE: 29-Oct-1999
CLASSIFICATION: <Unknown>
                      ATTORNEY/AGENT INFORMATION: NAME: Apple, Randolph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Andrews, William H.
TITLE OF INVENTION: No. 6309867el Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
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NAME: Apple, REGISTRATION
                                                                                           APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: Two Embarcadero Center,
                                                         APPLICATION NUMBER: US 08/724,643 FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Townsend and Townsend and Crew LLP
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Harley, Calvin
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Chapman, Karen B.
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                                                                                                                                                                                                                                                                                                                                                       PatentIn Release #1.0, Version #1.30
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Randolph T.
NUMBER: 36,429
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SEQUENCE DESCRIPTION: SEQ ID NO: 171:
US-09-430-323-171
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US-09-402-181B-291
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GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cech,
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TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/402,181B FILING DATE: 29-Sep-1997 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center; Eighth Floor CITY: San Francisco
                                      APPLICATION NUMBER: US 08/911,312 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: California
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                                                                             APPLICATION NUMBER: US 08/854,050 FILING DATE: 09-MAY-1997
                                                                                                                    FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
                                                                                                                                                                         APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
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  APPLICATION NUMBER: US 08/912,951 FILING DATE: 14-AUG-1997
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Nakamura, Toru
Chapman, Karen B.
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77.8%;
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Pred. No. 9
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RESULT 16
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                                               COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/721,456
FILING DATE: 22-No. 6617110-2000
CLASSIFICATION: <a href="https://doi.org/10.1002/00.00">https://doi.org/10.1002/00.00</a>
CLASSIFICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         41.0%;
Local Similarity 77.8%;
es 7; Conservar:...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Morin, Gregg B.
Harley, Calvin B.
Andarews, William H.
Andarews, William H.
NUMBER OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   telomerase core protein 1 (7 SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RMATION FOR SEQ ID NO: 291 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: WO PCT/US97/17885 FILING DATE: 01-OCT-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 LVMGVPFYG 11
                                                                                                                                                                                                                                                                                                                                                                              STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Peptide LOCATION: 1..15
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                    CITY: San Francisco
                      APPLICATION NUMBER: US 08/844,419 FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                                                                        ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/915,503 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/09721456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Thomas R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            291:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 34; DB
Pred: No. 9.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "motif D peptide from human
ein 1 (TCP1)"
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08/846,017
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4 LVRGVPEYG 12

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Query Match
Best Local Similarity
The 7; Conserve
                                                                                                                US-09-721-456-291
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                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                            OTHER INFORMATION: /note= "motif D peptide from human telomerase core protein 1 (TCP1)"
SEQUENCE DESCRIPTION: SEQ ID NO: 291:
                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                FEATURE:
3 LVMGVPFYG 11
                                                                                                                                                                      NAME/KEY: Peptide LOCATION: 1..15 OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                            NAME/KEY:
                                                                                                                                                                                                                                                                    STRANDEDNESS: <Unknown>
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: WO PCT/US97/17885 FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/851,843 FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                              LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/911,312 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/854,050 FILING DATE: 09-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 25-APR-1997
                                      Conservative
                                                        41.0%;
77.8%;
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                                                      Score 34; I
Pred. No. 9.
                                      Mismatches
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                                                      DB
9.3;
                                                                        2
                                                                      Length 15
                                      Indels
                                      0
                                      Gaps
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US-09-766-253-171 Sequence 171, Applicat Patent No. 6808880 GENERAL INFORMATION: Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: No. 6808880el To ZIP; 94111 COMPUTER READABLE FORM: CORRESPONDENCE ADDRESS: MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 ADDRESSEE: Townsend and STREET: Two Embarcadero COUNTRY: United States STATE: California CITY: San Francisco Application US/09766253 Cech, Thomas R. Lingner, Joachim of America Townsend and Crew Center, 8th Floor Telomerase

LLP

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NAME/KEY: Peptide;
; LOCATION: 1..15
; OTHER INFORMATION: /note= "motif D peptide from human telomerase core protein 1 (TCP1)"
; SEQUENCE DESCRIPTION: SEQ ID NO: 171:
US-09-766-253-171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-054-295-171
; Sequence 171, Applicat:
; Patent No. 6921664
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 171:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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APPLICATION NUMBER: 08/846,017
FILING DATE: 1997-04-25
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                        ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                            Andrews, William H. TITLE OF INVENTION: No. 6921664el Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 LVMGVPFYG 11
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                                                                                                                                                                                                                                                                                       STREET: Two Embarcadero Center,
CITY: San Francisco
STATE: California
                                                                          APPLICATION NUMBER: US/10/054,295
FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/766,253
FILING DATE: 19-Jan-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002920US
                  APPLICATION NUMBER: 08/854,050 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                         COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Townsend and
APPLICATION NUMBER: US 08/846,017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/10054295
                                                                                                                                                                                                                                                                                                                                                                                            OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Morin, Gregg B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ingner, Joachim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Karen B.
                                                                                                                                                                                                                                                                                                                                                                                            225
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                                                                                                                                                                                                                                                                                                                                 Townsend and Crew LLP Center, 8th Floor
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APPLICANT: LINGNER, JOACHIM
APPLICANT: LINGNER, JOACHIM
APPLICANT: CHAPMAN, KAKEN B.
APPLICANT: CHAPMAN, KAKEN B.
APPLICANT: CHAPMAN, KAKEN B.
APPLICANT: MORIN, GREGG B.
APPLICANT: MORIN, GREGG B.
APPLICANT: ANDREWS, WILLIAM H.
ITILE OF INVENTION: GENE FOR HUMAN TELOMERASE REVERSE TRANSCRIPTASE AND TITLE OF INVENTION: TELOMERASE VARIANTS
FILE REFERENCE: 018/062
FILE REFERENCE: 018/062
CURRENT APPLICATION NUMBER: US/09/438,486A
CURRENT FILING DATE: 1997-01-12
PRIOR APPLICATION NUMBER: 08/851,843
PRIOR APPLICATION NUMBER: 08/864,017
PRIOR APPLICATION NUMBER: 08/844,419
PRIOR APPLICATION NUMBER: 08/844,419
PRIOR APPLICATION NUMBER: 08/844,419
PRIOR APPLICATION NUMBER: 08/724,643
PRIOR APPLICATION NUMBER: 08/724,643
PRIOR FILING DATE: 1997-04-18
PRIOR FILING DATE: 1996-10-01
NUMBER OF SEQ ID NOS: 223
SOFTWARE: PATENTIN Ver. 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; telomerase core protein 1 (TCP1)"; SEQUENCE DESCRIPTION: SEQ ID NO: 171:
US-10-054-295-171
; OTHER INFORMATION: Description of Artificial Sequence: Recombinant; OTHER INFORMATION: amino acid sequence US-09-438-486A-171
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 171:
SEQUENCE CHARACTERISTICS:
                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                     ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION:
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: <Unknown>
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 15 amino acids
TYPE: amino acid
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1..15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41.0%;
77.8%;
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Pred. No. 9.3;
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Query Match

41.0%;

Score 34;

DB 2;

Length 15;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   밁
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                                                                                                                                          US-09-657-757-38
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SOFTWARE: Pate
SEQ ID NO 38
FENGTH: 16
                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US/09/171,705
PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 78
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 38
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Best Local Similarity
Matches 6; Conserv
                                                                                                     Query Match
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Sequence 38, App-
Sequence 38, App-
No. 618420/
                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 6881824
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                               APPLICANT: VERHEIJDEN, GILBERTUS F.M.
TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR USE IN ANTIGEN SPECIFIC TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY
FILE REFERENCE: 0/96198 US
CURRENT APPLICATION NUMBER: US/09/657,757
CURRENT FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BOOTS, ANNA M.H.
APPLICANT: VERHEIJDEN, GILBERTUS F.M.
TITLE OF INVENTION: NOVEL PEPTIDES SUITABLE FOR USE IN ANTIGEN SPECIFIC TITLE OF INVENTION: IMMUNOSUPPRESSIVE THERAPY
FILE REFERENCE: 0/96198 US
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: BOOTS, ANNA M.H. APPLICANT: VERHEIJDEN, GIL
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CURRENT FILING DATE: 1999-02-09
NUMBER OF SEQ ID NOS: 78
                                                                                                                                                     FEATURE:
OTHER INFORMATION: Description
OTHER INFORMATION: SEQUENCE OF
                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                             ENGTH:
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                                  2 KLVMGVP 8
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                                                                                      Similarity
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KLVMGIP 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KLVMGVP 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LVRGVPEYG 12
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                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11
                                                                                    41.0%;
85.7%;
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85.7%;
                                                                  Score 34; DB 2
Pred. No. 10;
1; Mismatches
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Pred. No. 10;
                                                                                                                                                        of Artificial Sequence: DERIVED FROM HUMAN CARTILAGE (HC) -39 PROTEIN
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                                                                                                   Length 16;
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                                                                    Indels
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US-10-038-972A-4
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US-08-444-818-420
                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: J. BARTLETT
TITLE OF INVENTION: AAV VECTORS AND METHODS
FILE REFERENCE: 28335/36996US
CURRENT APPLICATION NUMBER: US/10/038,972A
CURRENT FILING DATE: 2002-01-04
                                               SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 11
                                                                                                                                                                                                                                                                     Sequence 4, Applica
Patent No. 6962815
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                         PRIOR APPLICATION NUMBER: US 60/260,124 PRIOR FILING DATE: 2001-01-05 NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 14-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
ORGANISM: Artificial Sequence FEATURE:
                                    TYPE: PRT
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ZIP: 94608-2916
COMPUTER READABLE FORM:
COMPUTER READABLE Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Harbin, Alisa A.
REGISTATION NUMBER: 33,895
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (508)359-3876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/444,818
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APPLICANT: Rutter, William J.
TITLE OF INVENTION: NAMBY Dia
NUMBER OF SEQUENCES: 777
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CITY: Emeryville
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Pred. No. 4.6e+05
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; Patent No. 5460961
; Patent No. 5460961
APPLICANT: DEBY, CAROL; PINCEMAIL, JOEL; BOLLEN, ALEX
TITLE OF INVENTION: HUMAN MYELOPEROXIDASE AND ITS
; THERAPEUTIC APPLICATION
INVERER OF SEQUENCES: 13
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/641,678
FILING DATE: 16-JAN-1991
; PRIOR APPLICATION DATA:
; APPLICATION UMBER: 460,931
; PRIOR APPLICATION DATA:
; APPLICATION DATA:
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US-08-968-542C-22
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Matches
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Best Local Similarity 83.3%;
Matches 5; Conservative
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APPLICANT: Myers, et al.
TITLE OF INVENTION: dull1 Codes For A No. 5981728el Starch
TITLE OF INVENTION: Synthase
NUMBER OF SEQUENCES: 35
                                                                                                                                                                                                                    COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh
SOFTWARE: Microsoft Word 6.0.1 for Macintosh
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/968,542C
FILING DATE: No. 5981728ember 12, 1997
CLASSIFICATION BOO
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 floppy disk
ATTORNEY AGENT INFORMATION:
NAME: Benjamin Aaron Adler, Ph.D.,
REGISTRATION INVMBER: 35,423
REFERENCE/DOCKET NUMBER: D6036
TELECOMMUNICATION INFORMATION:
TELEPHONE: (713) 777-2321
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STREET: 8011 Candle Lane
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Search completed: January 26, 2006, 08:07:02 Job time : 17.1408 secs
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                                                                                                                            Query Match
Best Local S
                                                                                                                Matches
                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                   DESCRIPTION: amino acid
HYPOTHETICAL: no
ANTI-SENSE: no
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                             y Match
Local Similarity 62.5%;
hes 5; Conservative
                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                    1 DKTAMGAD 8
                                                                                                                                                                                                                                                                    1: 10 amino acid residues amino acid
                                                        DKSIVGVP 8
                                                                                                                                                                                                                                                          linear
                                                                                                                Score 30; DB Pred. No. 33; 2; Mismatches
                                                                                                                                            DB 1; Length 10;
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Run 3

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Result
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Maximum
Listing
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Perfect score:
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score greater than
and is derived by a
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83
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
      DKLVMGVPFYGRAXSIE
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Copyright (c) 1993 - 2006 Compugen
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US-09-791-393-111
US-09-791-379-311
US-09-791-377-324
US-10-264-309-277
US-10-264-309-277
US-10-264-309-277
US-10-264-309-278
US-10-264-309-278
US-10-371-525-273
US-10-371-645-273
US-10-371-645-273
US-10-371-645-273
US-10-371-645-273
US-10-371-645-273
US-10-371-650-273
US-10-371-651-165-146
US-10-388-084-1
US-09-766-253-171
US-09-748-486-171
US-09-438-486-171
US-10-054-295-171
US-10-054-611-171
US-10-054-71-172-291
US-10-877-022-291
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10, Appl
324, App
321, App
311, App
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 291, App
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425, App
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1653, App
1762, App
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1764, App
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1765, App
1766, App
1767, App
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ALIGNMENTS

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RESULT 2
US-09-791-378-324
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CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: US/09/292,225
PRIOR FILING DATE: 1999-04-15
PRIOR APPLICATION NUMBER: 60/098,909
PRIOR FILING DATE: 1998-09-02
PRIOR FILING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/085,295
PRIOR APPLICATION NUMBER: 60/085,295
PRIOR APPLICATION NUMBER: 60/085,295
PRIOR FILING DATE: 1998-05-13
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                                                                                                                                                                                           SOFTWARE: PatentIn version 3.0 SEQ ID NO 324
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 324, Application US/09791378 Patent No. US20020142303A1
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Best Local Similarity
                                Best Local Similarity
Matches 7; Conserv
                                                                       Query Match
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                                                                                                                                                                                                                                                                                         APPLICANT: Parekh, Rajesh
TITLE OF INVENTION: PROTEINS, GENES AND THEIR USE FOR DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: SCHIZOPHRENIA
FILE REFERENCE: 9195-061-999
CURRENT APPLICATION NUMBER: US/09/791,378
CURRENT FILING DATE: 2001-02-23
                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 09/750,395
PRIOR FILING DATE: 2000-12-28
NUMBER OF SEQ ID NOS: 677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PROTEINS TITLE OF INVENTION: AND USES THEREOF FILE REFERENCE: AL-2-C3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: McCall, Catherine A. APPLICANT: Hunter, Shirley Wu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.0
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PRIOR FILING DATE: 1998-04-17
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                                                                                                                                LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION:
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TYPE: PRT
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3 LVMGVPFYGR 12
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Weber, Eric R.
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ilarity 100.0%;
Conservative (
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                                                       49.4%;
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                                                       Score 41;
Pred. No.
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2.1;
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2.8e-07;
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APPLICANT: White, Frost
APPLICANT: Williams, Stephen A.
TITLE OF INVENTION: Nucleic Acid Molecules, Polyf
TITLE OF INVENTION: Nucleic Acid Molecules, Polyf
TITLE OF INVENTION: Uses Therefor, Including Dis
TITLE OF INVENTION: Alzheimer's Disease
FILE REFERENCE: 2572-1-001 N2
CURRENT APPLICATION NUMBER: US(19/826,290)
CURRENT FILING DATE: 2001-04-30
PRIOR APPLICATION NUMBER: US 60/194,504
PRIOR APPLICATION NUMBER: US 60/253,647
PRIOR FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 492
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Publication No. US20
GENERAL INFORMATION:
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Matches
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LENGTH: 10
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 111 LENGTH: 10
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                                                            EARLIER FILING DATE: 2000-12-12
NUMBER OF SEQ ID NOS: 308
                                                                                                  CURRENT APPLICATION NUMBER: US/09/791,393
CURRENT FILING DATE: 2002-01-02
EARLIER APPLICATION NUMBER: GB 0004412.3
EARLIER FILING DATE: 2000-02-24
EARLIER APPLICATION NUMBER: GB 0030050.9
EARLIER FILING DATE: 2000-12-08
EARLIER APPLICATION NUMBER: US 60/254,830
                                                                                                                                                                                                                                                     TITLE OF INVENTION: Proteins, Genes and Their Use for TITLE OF INVENTION: Diagnosis and Treatment of Bipolar Affective Disorder (BAD) TITLE OF INVENTION: and Unipolar Depression FILE REFERENCE: 2543-1-001 N1
                                                                                                                                                                                                                                                                                                                                  APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Rohlff, Christian
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7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09791393
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David M.
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B. Michael
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US-09-791-377-324
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; ORGANISM: homo sapien
US-09-791-393-111
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                                                                                    Sequence 324, Application US/09791377

Publication No. US20040110938A1

GENERAL INFORMATION:
APPLICANT: Parekh, Rajesh
TITLE OF INVENTION: PROTEINS, GENES AND THEIR USE FOR DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: SCHIZOPHRENIA
FILE REFERENCE: 9195-060-999
CURRENT FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: US/09/791,377
CURRENT FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 09/750,395
PRIOR APPLICATION NUMBER: 09/750,395
PRIOR TILING DATE: 2000-12-28
NUMBER OF SEQ ID NOS: 677
SOFTWARE: PATERING VERSION 3.0

SEQ ID NO 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 308
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 111
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Best Local Similarity 70.0%;
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Best Local Similarity 70.0%;
Matches 7; Conservative 2
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APPLICANT: Tygon, Kerry Louise
TITLE OF INVENTION: Proteins, Genes and Their Use for
TITLE OF INVENTION: Diagnosis and Treatment of Bipolar Affective Disorder (BAD)
TITLE OF INVENTION: and Unipolar Depression
FILE REFERENCE: 2543-1-01 N2
CURRENT APPLICATION NUMBER: US/09/791,389
CURRENT FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: GB 0004412.3
PRIOR APPLICATION NUMBER: GB 003050.9
PRIOR APPLICATION NUMBER: US 60/254,830
PRIOR PILING DATE: 2000-12-08
PRIOR PILING DATE: 2000-12-12
LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
-09-791-377-324
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APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Rohlff, Christian
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o. US20030032773A1
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Pred. No. 2.1;
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APPLICANT: WILLIAMS, STEPHEN A.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES, POLA
TITLE OF INVENTION: INCLUDING DIAGNOSIS AND TRE
FILE REFERENCE: POA-002.01
CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT FILING DATE: 2002-10-03
PRIOR APPLICATION NUMBER: 60/326,708
PRIOR APPLICATION NUMBER: 60/326,708
PRIOR FILING DATE: 2001-10-03
NUMBER: OF SEQ ID NOS: 491
SOFTWARE: Patentin Version 2.1
SEQ ID NO 277
LENGTH: 10
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                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                    Sequence 277, A Publication No.
                                                                                                                                                                                                                                                                                                                                                                       Query Match 49.4%;
Best Local Similarity 70.0%;
Matches 7; Conservative
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Matches
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APPLICANT:
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APPLICANT:
APPLICANT:
                                           APPLICANT
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                           APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
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                       KIMMEL, LIDA H.
PAREKH, RAJESH B.
POTTER, DAVID M.
ROHLFF, CHRISTIAN
SILBER, B. MICHAEL
SNYDER, PETER J.
SOARES, HOLLY D.
STIGER, THOMAS R.
SUNDERLAND, P. TREY
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LVMGIPTFGR 10
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SOARES, HOLLY D.
STIGER, THOMAS R.
SUNDERLAND, P. TREY
TOWNSEND, ROBERT R.
WHITE, W. FROST
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KIMMEL, LIDA H.
PAREKH, RAJESH B.
POTTER, DAVID M.
ROHLFF, CHRISTIAN
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FRIEDMAN, DAVID L.
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FRIEDMAN, DAVID L.
          TOWNSEND,
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                                                                                                                                                                                                                                   Application US/10264309
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5. US20040022794A1
                                                                                                                                                                                                                       US20050163789A9
ROBERT R.
FROST
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Pred. No. 2.1;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                       2; Mismatches
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Pred. No. 2.1;
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RESULT 10
US-10-264-309-278
iy Sequence 278, Application US/10264309
publication No. US20050163789A9
GENERAL INFORMATION:
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-264-309-278
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APPLICANT: WILLIAMS, STEPHEN A.
APPLICANT: WILLIAMS, STEPHEN A.
TITLE OF INVENTION: INCLUDING DIAGNOSIS AND TREATMENT OF ALZHEIMER'S DISEASE
FILE REFERENCE: POA-002.01
CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT FILING DATE: 2002-10-03
PRIOR APPLICATION NUMBER: 60/326,708
PRIOR APPLICATION NUMBER: 60/326,708
PRIOR PILING DATE: 2001-10-03
NUMBER OF SEO ID NOS: 491
SOPTWARE: Patentin Version 2.1
SEO ID NO 278
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US-10-264-309-278
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CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT FILING DATE: 2002-10-03
PRIOR APPLICATION NUMBER: 60/326,708
PRIOR APPLICATION NUMBER: 60/326,708
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 491
SOCTWARE: PATENTIN Version 2.1
SEQ ID NO 277
                                                                                                                                                                               Query Match
Best Local Similarity
7; Conserv
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APPLICANT:
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Matches 7; Conservative
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             POTTER,
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o. US20040022794A1
                                                                                                                                                                                               Conservative
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DAVID M.
CHRISTIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HOLLY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B. MICHAEL
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70.0%;
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Pred. No. 2.1;
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Pred. No. 4.8;
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APPLICANT: SUNDER, PETER J.
APPLICANT: SOARES, HOLLY D.
APPLICANT: SOARES, HOLLY D.
APPLICANT: SUNDERLAND, P. TRRY
APPLICANT: SUNDERLAND, P. TRRY
APPLICANT: TOWNSEND, ROBERT R.
APPLICANT: WHITE, W. FROST
APPLICANT: WHITE, W. FROST
APPLICANT: WHITE, W. FROST
APPLICANT: WILLIAMS, STEPHEN A.
TITLE OF INVENTION: INCLUDING DIAGNOSIS AND TREATMENT OF ALZHEIMER'S DISEASE
FILE REFERENCE: POA-002.01
FILE REFERENCE: POA-002.01
CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT APPLICATION NUMBER: 0031-0-03
PRIOR APPLICATION WHERE: 0032-10-03
PRIOR APPLICATION UNMBER: 6033
NUMBER OF SEQ ID NOS: 491
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US-09-017-743C-3
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LENGTH: 10
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Best Local
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             COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTMARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION NUMBER: US/09/017,743C
APPLICATION NUMBER: US/09/017,743C
FILING DATE: 03-Feb-1998
CLASSIFICATION ONTA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/590,298
ATTORNEY/AGENT INFORMATION:
NAME: Parent, Annette S.
REGISTRATION NUMBER: 42,058
REGISTRATION NUMBER: 42,058
                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: HLA Binding Peptides and Their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sette, Alessandro
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7; Conserv
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KIMMEL, LIDA H.
PAREKH, RAJESH B.
POTTER, DAVID M.
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                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                            STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: San Francisco
REFERENCE/DOCKET NUMBER: 018623-008050US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ROHLFF, CHRISTIAN
SILBER, B. MICHAEL
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Pred. No. 4.8;
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RESULT 13
US-10-371-069-273
Sequence 273, App
Publication No. L
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CURRENT FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: US 09/311,784
PRIOR FILING DATE: 1999-05-13
PRIOR APPLICATION NUMBER: US 60/085,751
PRIOR FILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 463
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 273
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Best Local Similarity
Marches 5; Conserve
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; OTHER INFORMATION: HCV 1378 (peptide 29.0035)
US-10-371-525-273
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US-10-371-525-273
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                                                                                    GENERAL INFORMATION:
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Best Local
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APPLICANT: EPIMMUNE Inc.
APPLICANT: Fikes, John D.
APPLICANT: Hermanson, Gary G.
APPLICANT: Sette, Alessandro
APPLICANT: Ishioka, Glenn Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Expression Vectors for Stimulating an TITLE OF INVENTION: Immune Response and Methods of Using the FILE REFERENCE: 39963-20022.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Fikes, John D.
APPLICANT: Hermanson, Gary G.
APPLICANT: Sette, Alessandro
APPLICANT: Ishioka, Glenn Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                     7 VPFYGRA 13
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-576-0300
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                                                                                                                    Application US/10371069
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                                                                                                    US20030216342A1
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71.4%;
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Pred. No. 1.7e+06;
Pred. No. 1.7e+06;
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Pred. No. 1
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Best Local Similarity
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; OTHER INFORMATION: HCV 1378 (peptide 29.0035)
US-10-371-069-273
                                                                                                                                                US-10-371-645-273
                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/078,904
PRIOR ETLING DATE: 1998-05-13
PRIOR APPLICATION NUMBER: US 60/085,751
PRIOR FILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 463
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 273
                                                                        Matches
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                                                                                                         Query Match
                                                                                          Best Local
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PRIOR FILING DATE: 1998-05-13
PRIOR APPLICATION NUMBER: US 60/085, 751
PRIOR FILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 463
                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ishioka, Glenn Y.
APPLICANT: Livingston, Brian
APPLICANT: Chesnut, Robert W.
APPLICANT: Epimmune Inc.
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: Expression Vectors for Stimulating an TITLE OF INVENTION: Immune Response and Methods of Using FILE REFERENCE: 39963-20022.11
CURRENT APPLICATION NUMBER: US/10/371,645
CURRENT FILING DATE: 2003-66-20
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APPLICANT: Epimmune Inc.
TITLE OF INVENTION: Expression Vectors for Stimulating an TITLE OF INVENTION: Immune Response and Methods of Using the FILE REFERENCE: 39963-20022.10
CURRENT APPLICATION NUMBER: US/10/371,069
CURRENT FILING DATE: 2003-02-21
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                  OTHER INFORMATION: HCV 1378 (peptide 29.0035)
                                                                                                                                                                                                                                            ENGTH:
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                                   7 VPFYGRA 13
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                                                                                          Similarity
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                                                                      5; Conservative
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                                                                                        42.2%;
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71.4%;
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Pred. No. 1.7e+06
2; Mismatches
                                                                                        Score 35; DB 4;
Pred. No. 1.7e+06;
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US-10-371-260-273

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                                                                                                              ; ORGANISM: hepatitis C virus US-10-651-165-146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-651-165-146
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                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/EP94/03555
PRIOR FILING DATE: 1994-10-28
PRIOR APPLICATION NUMBER: EP 93402718.6
PRIOR FILING DATE: 1993-11-04
NUMBER OF SEQ ID NOS: 286
SOFTWARE: Patentin version 3.1
SEQ ID NO 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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Best Local
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                                    Best Local Similarity
Matches 5; Conserv
                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Expression Vectors for Stimulating an TITLE OF INVENTION: Immune Response and Methods of Using the FILE REFERENCE: 39963-20022.13 CURRENT APPLICATION NUMBER: US/10/371,260 CURRENT FILING DATE: 2003-02-21
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                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/651,165
CURRENT FILING DATE: 2003-09-02
PRIOR APPLICATION NUMBER: US/08/974,690C
PRIOR FILING DATE: 1997-11-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: LEROUX-ROELS, Geert APPLICANT: DELEYS, Robert APPLICANT: MAERTENS, Geert
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APPLICANT: Fikes, John
APPLICANT: Hermanson, (
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SOFTWARE: FastSEQ for Windows Version 3.0
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ORGANISM: Artificial Sequence
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7 VPFYGRA 13
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Hermanson, Gary G.
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Ishioka, Glenn Y.
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                                      Conservative
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                                                          42.2%;
71.4%;
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                                                        Score 35; DB 4;
Pred. No. 1.7e+06;
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Pred. No. 1.
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US-09-843-676-171
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GENERAL INFORMATION:
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LENGTH: 21
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/888,084
CURRENT FILING DATE: 2004-07-09
PRIOR APPLICATION NUMBER: 60/486,581
PRIOR FILING DATE: 2003-07-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 1
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Aebersold, Ruedi APPLICANT: Zhang, Ruedi
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ORGANISM: Artificial Sequence
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                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/843,676
FILING DATE: 26-Apr-2001
CLASSIFICATION S56
PRIOR APPLICATION NUMBER: US/08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US/08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
TILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
                                                                                                                                                                                                                                                                                                                                                                                                                                                Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Greeg B.
Morin, Greeg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: No. US20020164786A1el Telomerase
NUMBER OF SEQUENCES: 225
CORRESPONDENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
ADDRESSEE: Townsend and Townsend and Crew LLP
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STATE: California
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18-APR-1997
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IOCATION: 1.15
; OTHER INFORMATION: /note= "motif D peptide from human;
; telomerage core protein 1 (TCP1)"
; SEQUENCE DESCRIPTION: SEQ ID NO: 171:
US-09-843-676-171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 19
US-09-766-253-171
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          APPLICATION NUMBER: 08/846,017
FILING DATE: 1997-04-25
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: APPLe, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002920US
                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/766,253
FILING DATE: 19-Jan;2001
CLASSIFICATION: <UKnown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Morin, Gregg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: No. US20020187471A1el Telomerase
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LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                             CITY: San Francisco
STATE: California
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REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor
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Nakamura, Toru
Chapman, Karen B.
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Pred. No. 58;
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US-09-438-486-171
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Matches 7; Conserv
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                                       CLASSIFICATION: 536
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/724,643
                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                             CLASSIFICATION: 536 PRIOR APPLICATION DATA:
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APPLICANT: Chapman, Karen B.
APPLICANT: Morin, Gregg B.
APPLICANT: Harley, Calvin
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                                                                                                                                                         APPLICATION NUMBER: US 08/846,017 FILING DATE: 25-APR-1997 CLASSIFICATION: 536
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CLASSIFICATION:
                                APPLICATION NUMBER: FILING DATE: 01-00
                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/438,486
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SEQUENCE DESCRIPTION: SEQ ID NO: 171:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: peptide
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California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T: Andrews, William H. INVENTION: No. US20030009019A1el Telomerase
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OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
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TELEFAX: (415) 576-0300
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b. US20030009019A1
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INFORMATION:
                              01-OCT-1996
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77.8%;
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; LOCATION: 1..15
; OTHER INFORMATION: /note= "motif D peptide from human;
OTHER INFORMATION: telomerase core protein 1 (TCP1)"
US-09-438-486-171
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Best Local
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INFORMATION FOR SEQ ID NO: 171:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200
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                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
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                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/053,758
FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Andrews, William H.
TITLE OF INVENTION: No. US20030032075Alel Telomerase
NUMBER OF SEQUENCES: 225
                     ATTORNEY/AGENT INFORMATION: NAME: Apple, Randolph
                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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Similarity 77.8%;
7; Conservation
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Embarcadero Center, 8th Floor
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NAME: Apple,
REGISTRATION
                                                       APPLICATION NUMBER: US 08/724,643 FILING DATE: 01-OCT-1996
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Harley, Calvin
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Chapman, Karen B.
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 NUMBER: 36,429
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             Randolph T.
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Pred. No. 58;
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US-10-054-295-171
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INFORMATION FOR SEQ ID NO: 171:
SEQUENCE CHARACTERISTICS:
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Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Morin, Gregg B.
Harley, Calvin
Andrews, William H.
ANDREWS, William H.
FITLE OF INVENTION: No. US20030044953A1el Telomerase
NUMBER OF SEQUENCES: 225
CORRESPONDENCE ADDRESS:
                                APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/854,050
FILING DATE: <Unknown>
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OTHER INFORMATION: /note= "motif D peptide from human telomerase core protein 1 (TCP1)"
SEQUENCE DESCRIPTION: SEQ ID NO: 171:
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TELEPHONE: (415) 576-0200
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                   TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/10/054,295 FILING DATE: 18-Jan-2002 CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                              APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 015389-002930US
TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 15 amino acids
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o. US20030044953A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide
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77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thomas R.
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Pred. No. 58;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Townsend and Crew LLP
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RESULT 23
US-10-054-611-171
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Best Local
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TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 171: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 171: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION NUMBER: US/10/054,611
FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/854,050

FILING DATE: <Unknown:
APPLICATION NUMBER: US 08/846,017

FILING DATE: 25-APR-1997

APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997

APPLICATION NUMBER: US 08/724,643

FILING DATE: 01-OCT-1996

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                    NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Andrews, William H.
TITLE OF INVENTION: No. US20030059787A1el Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Peptide
LOCATION: 1..15
LOCHER INFORMATION: /note= "motif D peptide from human telomerase core protein 1 (TCP1)"
SEQUENCE DESCRIPTION: SEQ ID NO: 171:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: San Francisco
STATE: California
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LVRGVPEYG 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 15 amino acids
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Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin
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; telomerase core protein 1 (TCP1)" ; SEQUENCE DESCRIPTION: SEQ ID NO: 171: US-10-054-611-171
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US-10-325-810-291
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US-10-282-960-70
                                                                                                                                                                                          Sequence 291, Application US/1032 Publication No. US20030204069A1 GENERAL INFORMATION:

APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Chen, Si-Yi
APPLICANT: Zhaoyang, You
APPLICANT: Schroers, Roland
TITLE OF INVENTION: Human Telomerase Reverse Transcriptase as a
TITLE OF INVENTION: Associated Antigen
FILE REFERENCE: P02193US1
CURRENT APPLICATION NUMBER: US/10/282,960
CURRENT FILING DATE: 2002-10-29
CURRENT FILING DATE: 2002-10-29
                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin
SEQ ID NO 70
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                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/345,012
PRIOR FILING DATE: 2001-10-29
NUMBER OF SEQ ID NOS: 100
SOFTWARE: Parcer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 15
TYPE: PRT
ORGANISM: Human
                                                           Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
           CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
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7; Conservative
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LOCATION: 1.15
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/10282960
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5. US20030204069A1
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77.8%;
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Pred. No. 58;
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Pred. No. 58;
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Search completed: January 26, 2006, 08:38:37 Job time : 56.25 secs
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Best Local Similarity 77.8%;
Matches 7; Conservative
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TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 291:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/854,050
APPLICATION NUMBER: US 08/911,312
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
APPLICATION NUMBER: US 08/915,503
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY, AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: /note= "motif D peptide from human telomerase core protein 1 (TCP1)" SEQUENCE DESCRIPTION: SEQ ID NO: 291:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/325,810
FILING DATE: 20-Dec-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/402,181
FILING DATE: 29-Sep-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
                                                                                                                                          3 LVMGVPFYG 11
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LVRGVPEYG 12
                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: Peptide LOCATION: 1.15
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: <Unknown>
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 15 amino acids TYPE: amino acid
                                                                                                                                                                                     Score 34; DB 4; Length 15; Pred. No. 58; 0; Mismatches 2; Indels
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Maximum Match 100%
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192.434 Million cell updates/sec
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Copyright (c) 1993 - 2006 Compugen Ltd
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                                 flavodoxin A Azo
protein phosphatas
Ig heavy chain DJ
T-cell receptor al
unknown protein en
sialokinin II - ye
sialokinin II - yel
LuxC protein - Pho
tetrahydroberberin
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Ig heavy chain CRD
chlorophyll a/b-bi
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unidentified 5.4/3
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hydrogenase (EC 1.
T cell receptor al
hypothetical TCL3/
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              B61033
A24867
ECLQ2M
A26120
S07201
S07201
A61033
C39109
S07436
A32486
A32486
                   PH0809
DRUFPD
JS0647
PT0239
PH1794
A45590
S43360
A28814
PT0332
I49039
A24589
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140665
S26747
PH1789
B31769
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JN0390
PA0015
B39111
S48685
                                                                                                                                       PA0014
PN0173
                                                                                                                                                                                                                                                                                                           B12055
A27719
S09088
F64121
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S53440
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S51734
S50022
S46205
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C49048
A41170
B36727
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A56970
S04586
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A40693
S21152
PA0013
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hypothetical prote hypothetical BCR/A fibrinopeptide B - tachykinin I - mig tachykinin II - mig tachykinin II - mig cytochrome P450 1A ranatachykinin II - s cyliorhinin II - mi 6-phosphofructokin uperolein - frog yhysalaemin - frog ranatachykinin A -

ranatachykinin A hypothetical 1.2K tachykinin - Afric

beta protein - rat seed protein ws-4 bradykinin-like pe

histamine-releasin

glyceraldehyde-3-p trypsin (EC 3.4.21 proteasome chain 7 hypothetical prote

hypothetical prote glutathione-bindin

metalloproteinase

photosystem II 6.1 cytochrome c554 proteasome chain 5 brain-derived neur T-cell receptor al trypsin-like prote comosain (EC 3.4.2

tryptophyllin-rela photosystem II oxy protein QA600027 -GLYMA1 - soybean (NADH2 dehydrogenas T-cell receptor al

transgelin - sheep

ig heavy chain CDR
T cell receptor al
beta-pigment dispe
Ig mu chain V regi
Ig kappa chain CRD
T cell receptor be

T-cell receptor be photosystem II chl hypothetical prote Amino terminal of

ilvBN leader pepti Ig heavy chain J T cell receptor al T-cell receptor de T-cell receptor al pigment-dispersing pheromonotropin

seed storage prote seed storage prote T-cell receptor al ilvBN leader pepti

seed storage prote Ig heavy chain V rextension protein phe operon leader

photosystem II ext distal retinal-pig proteasome endopep 37K adherens junct Ig H chain V-D-J r Ig heavy chain DJ

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R;Kovacs, K.L.; Tigyi, G.; Thanh, L.T.; Lakatos, S.; Kiss, Z.; Bagyinka, C. J. Biol. Chem. 266, 947-951, 1915
A;Title: Structural rearrangements in active and inactive forms of hydrogenase A;Reference number: A39089; MUID:91093297; PMID:1845998
A;Accession: B33089
                                                                                                                                                                                                                                                hydrogenase (EC 1.18.99.1) 34K chain - Thiocapsa roseopersicina (fragment) C;Species: Thiocapsa roseopersicina C;Date: 27-Nov-1991 #sequence_revision 27-Nov-1991 #text_change 23-Jun-1993 C;Accession: B39089
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      glutamate dehydrogenase (EC 1.4.1.2) - bluefin tuna (tentative sequence) (f C;Species: Thunnus thynnus (bluefin tuna)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 05-Oct-2004
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A; Residues: 1-7 <BAC>
A; Cross-references: UNIPARC: UPI000017C6D9;
C; Keywords: immunoglobulin
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A;Title: Two murine natural polyreactive autoantibodies are encod. A;Reference number: A33932; MUID:89282823; PMID:2499887
A;Accession: E33932
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C;Species: Mus musculus (house mouse)
C;Date: 09-Mar-1990 #sequence_revision 09-Mar-1990 #text_change 16-Aug-1996
C;Accession: E33992
                                                                                       A; Molecule type: protein A; Residues: 1-20 < KOV>
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R;Veronese, F.M.; Bevilacqua, R.; Boccu, E.;
                                                                                                                         A; Status: preliminary
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                                                                    Cross-references: UNIPARC:UPI000017AB7F
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        Query Match
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35.7%;
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        30.1%;
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Pred. No. 3.3e+02;
3; Mismatches 6
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Pred. No. 2.8e+05
        Score
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                         A;Title:
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RESULT 4
PH1763
C1 receptor alpha chain V region (clone 1V alpha 23-2) - human (fragment C;Species: Homo sapiens (man)
C;Species: 16-Ull-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
R;Porcelli, S.; Yockey, C.E.; Brenner, M.B.; Balk, J. Exp. Med. 178, 1-16, 1993
A;Title: Analysis of T cell antigen receptor (TCR)
                                                                                                       T cell receptor alpha chain V region (clone 5PBL V alpha 24-3) C;Species: Homo sapiens (man) C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change C;Accession: PH1820
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C;Keywords: fusion protein
F;11-5/Region: TC13 proto-oncogene derived
F;10-16/Region: T-cell receptor delta chain derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical TCL3/TCRD (T-cell receptor delta chain) mutant fusion protein - C;Species: Homo sapiens (man)
C;Date: 28-Jan-2000 #sequence_revision 28-Jan-2000 #text_change 09-Jul-2004
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C;Comment: This sequence is the hypothetical translation of a chimeric translocation mut
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A; Residues: 1-16 < ZUT>
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A;Title: The t(10;14)(q24;q11) of T-cell acute lymphoblastic leukemia juxtaposes that a serious control of the co
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J. Exp. Med. 178, 1-16, 1993
A;Title: Analysis of T cell antigen receptor
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A; Residues: 1-14 < POR>
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66.7%;
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Pred. No. 6.4e+02;
1; Mismatches 1
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Pred. No. 7.
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2; Mismatches 0;
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by human peripheral
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A;Reference number:
A;Accession: PH1820
A;Status
                                                                             RESULT 9
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Theor. Appl. Genet. 86, 935-942, 1993
A;Title: A rice protein library; a data-file
A;Reference number: PQ0696
A;Accession: PQ0730
A;Accession: PQ0730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: 146652
R;Yang, Y.G.; Ohta, S.; Yamada, S.; Shimizu, M.; Takagaki, Y.
J. Immunol. 155, 1981-1993, 1995
A;Title: Diversity of T cell receptor delta-chain cDNA in the A;Reference number: 146623; MUID:95363165; PMID:7636249
A;Accession: 146652
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I46652
NADH2 dehydrogenase (EC 1.6.99.3) 14K chain - fava bean mitochondrion (fragn N;Alternate names: complex I 14K chain; NADH-ubiquinone reductase 14K chain C;Species: mitochondrion Vicia faba (fava bean) C;Date: 03-May-1994 #sequence_revision 07-Oct-1994 #text_change 09-Jul-2004
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C;Accession: PQ0730
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A; Residues: 1-20 < YAN>
                                                                                                                                                                                                                                                                                                                A; Molecule type: protein A; Residues: 1-12 < KOM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unidentified 5.4/35K protein [imported] -
C;Species: Oryza sativa (rice)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T-cell receptor delta-chain J-delta-1 segment - pig (fragment)
C;Species: Sus scrofa domestica (domestic pig)
C;Species: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change
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A; Residues: 1-17 < POR>
                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:Q7M1U3; UNIPARC:UPI000017B124
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Best Local
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66.7%;
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Pred. No. 7.9e
1; Mismatches
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Pred. No. 8.2e+02;
0; Mismatches 1
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Pred. No. 9.3e+02;
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.9e+02;
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K; DAROCHE, J.; Bennett, J.; Falkowski, P.G.
Gene 95, 165-171, 1990
A;Title: Characterization of a cDNA encoding for the 28.5-kDa LHCII apoprotein A;Reference number: JW0040; MUID:91065528; PMID:2249775
A;Accession: PW0004
                                                                                                                             C; Superramıı
C; Keywords:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chlorophyll a/b-binding protein 24.5K - green alga N;Alternate names: photosystem II light-harvesting C;Species: Dunaliella tertiolecta C;Date: 04-Sep-1998 #sequence_revision 24-Nov-1999 C;Pate: 04-Sep-1998 #sequence_revision 24-Nov-1999 C;Accession: Dunnow
                                                                                                                     A;Cross-references: UNIPARC:UPI0000178174
C;Superfamily: chlorophyll a/b-binding protein
C;Keywords: chloroplast; grana; light-harvesting complex; membrane adhesion; membrane
                                                                                                                                                                                                                                        A; Molecule type: protein A; Residues: 1-15'<LAR>
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A;Title: Preferential utilization of specific immunoglobulin heavy chain div. A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0324
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C;Superfamily: NADH dehydrogenase
C;Keywords: electron transfer; mitochondrion;
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C;Comment: Complex I, mitochondrial
ranging from 5K to 75K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C; Keywords:
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A; Residues: 1-9 < YAM>
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Score 22; DB
Pred. No. 1.6e
0; Mismatches
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Pred. No. 1.3e+03;
1; Mismatches 1
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Pred. No. 2.8e+05
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1.6e+03;
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S57584
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A;Description: T cell receptor repertoire for a viral epitope in humans is diversified A;Reference number: S57494
A;Accession: S57584
                                                                                                                                                                                                                                   RESULT 14
PH1801
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PH1782
C:Species: Homo sapiens (man)
C:Species: 16-Ull-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
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A;Reference number: PH1754;
A;Accession: PH1801
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-17 <POR>
                                                                                                                                                            T cell receptor alpha chain V region (clone 3PBL V alpha 24-7) - C;Species: Homo sapiens (man) C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 1 C;Accession: PH1801
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                                                                                                         R;Porcelli, S.; Yockey, C.E.; Brenner, M.B.; Balk, S.P. .
J. Exp. Med. 178, 1-16, 1993
A;Title: Analysis of T cell antigen receptor (TCR) expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-16 < POR>
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A;Title: Analysis of T cell
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J. Exp. Med. 178, 1-16, 1993
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A; Residues: 1-15 <BUR>
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Pred. No. 1.7e
2; Mismatches
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Pred. No. 1.6e+03;
2; Mismatches 1
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C;Accession: S16929
R;Bagby, S.; Barker, P.D.; Hill, H.A.O.; Sanghera, G.S.; Dunbar, B.; Ashby, G.A.; Eady, Biochem. J. 277, 313-319, 1991
A;Title: Direct electrochemistry of two genetically distinct flavodoxins isolated from A;Reference number: S16929; MUID:91315397; PMID:1859358
A;Accession: S16929
Ig heavy chain DJ region (clone C439-111) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1304
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G. J. Exp. Med. 176, 1577-1581, 1992
                                                                                                           RESULT 17
PH1304
                                                                                                                                                                                                                                                                                                                                                                                                   R;Tamura, S.; Yasui, A.; Tsuiki, S.
Biochem. Biophys. Res. Commun. 163, 131-136, 1989
A;Title: Expression of rat protein phosphatase 2C (IA)
A;Reference number: A32917; MUID:89374212; PMID:2549985
A;Accession: A32917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein phosphatase - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 08-Dec-1989 #sequence_revision 08
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A;Residues: 1-17 <BAGS
A;Cross-references: UNIPROT:P23002; UNIPARC:UPI000012A8CA
A;Experimental source: strain MCD1155
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A;Residues: 1-18 <TAM>
A;Cross-references: UNIPARC:UPI000017C9A7
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;Date: 19-Mar-1997 #sequence_revision 11-Jun-1999
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Pred. No. 1.8e+03;
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Pred. No. 1.9e+03
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A;Title: Predominance of fetal type DJH joini A;Reference number: PH1302; MUID:93094761; PM A;Accession: PH1304 A;Molecule type: DNA A;Molecule type: DNA A;Cross-references: UNIPARC:UPI0000176934 C;Superfamily: immunoglobulin V region; immun C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                A;Experimental source:
C;Genetics:
A;Gene: Z2087
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                                                                                                                                                                                                                                                                                         A;Title: Genome sequence of enterohemorrhagic Escherichia A;Reference number: A85480; MUID:21074935; PMID:11206551 A;Accession: A85712
                                                                                                                                                                                                                                                                                                                                             iller, L.; Grotbeck, E.J.; Davis, Nature 409, 529-533, 2001
                                                                                                                                                                                                                                                                                                                                                           R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glassiller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta,
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S03504
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                                                                                                                                                                                                                                                        A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R; Winoto, A.; Mjolsness, S.; Hood, L. Nature 316, 832-836, 1985
A; Title: Genomic organization of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                         unknown protein encoded by prophage CP-9330 [imported] -
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A; Residues: 1-21 <WIN>
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C;Species Mus musculus (house mouse)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_chang
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Best Local S
Matches 4
                                                                                                                                                                                                ;Cross-references: UNIPROT:Q8X4E1; UNIPARC:UPI0000DDDFB; GB:AE005174; NID:g12515036;
;Experimental source: strain O157:H7, substrain EDL933
                                                                                                                                                                                                                                                                                                                                                                                                               Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                          Species: Escherichia col:
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Pred. No. 2e+C
1; Mismatches
                                                                                        Score 22; DB 2; Lo
Pred. No. 2.2e+03;
3; Mismatches 5;
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Pred. No. 2.2e+03;
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PMID:2993908
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2e+03;
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)imalanta, E.;
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homology
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K.; Apodaca,
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Query Match
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R;Ferri, S.R.; Meighen, E.A.
J. Biol. Chem. 266, 12852-12857, 1991
A;Title: A lux-specific myristoyl transferase in luminescent bacteria related to A;Reference number: A39853; MUID:91302295; PMID:2071574
A;Accession: B39853
                                                                                                                                                                                                   LuxC protein - Photobacterium phosphoreum (fragment)
C;Species: Photobacterium phosphoreum
C;Date: 30-Dec-1991 #sequence_revision 30-Dec-1991 #text_change 09-Jul-2004
                      A; Molecule type: DNA
A; Residues: 1-11 <FEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Accession: A49581
R;Champagne, D.E.; Ribeiro, J.M.
Proc. Natl. Acad. Sci. U.S.A. 91, 138-142,
A;Title: Sialokinin I and II: vasodilatory
A;Reference number: A49581; MUID:94105119;
A;Accession: A49581
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sialokinin I - yellow fever mosquito
C;Species: Aedes aegypti (yellow fever
C;Date: 06-Oct-1994 #sequence_revision
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Proc. Natl. Acad. Sci. U.S.A. 91, 138-142,
A;Title: Sialokinin I and II: vasodilatory
A;Reference number: A49581; MUID:94105119;
A;Accession: B49581
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(;Species: Aedes aegypti (yellow fever mosquito)
C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 16-Aug-2004
C;Date: 06-Oct-1994
                                                              A;Status: preliminary
                                                                                                                                                                                            C; Accession: B39853
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UNIPROT: P19841; UNIPARC: UPI000016FDC7; GB:M64224;
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66.7%;
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Pred. No. 1.5e+03;
0; Mismatches 2
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Pred. No. 1.
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18-Nov-1994
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y tachykinins
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tachykinins from
PMID:8278354
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1.5e+03;
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Similarity

25.3%;

Score 21; Pred. No.

DB 2; 1.7e+03;

Length 11;

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A;Reference number: A44957
A;Accession: G44957
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-14 <TAK>
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G44957
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submitted to JIPID, April 1993
A;Reference number: PS0206
A;Accession: PS0325
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B61620
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C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 11-Apr-1995
C;Accession: PS0325
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PS0325
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                                                                                                photosystem II oxygen-evolving complex protein 2 - common tobacco (cv. Samsun NN) (fragn C;Species: Nicotiana tabacum (common tobacco) C;Date: 03-Jun-1993 #sequence_revision 03-Jun-1993 #text_change 18-Jun-1993 C;Accession: G44957 R;Takeda, S.; Sato, F.; Ida, K.; Yamada, Y. Plant Cell Physiol. 31, 215-221, 1990 A;Title: Characterization of polypeptides that accumulate in cultured Nicotiana tabacum
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A;Title: Isolation, identification and synthesis of locustamyotropin III and IV, two add A;Reference number: A61620
A;Accession: B61620
A;Status: preliminary
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C;Keywords: amidated carboxyl end; neuropeptide
F;13/Modified site: amidated carboxyl end (Leu) #status experimental
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A;Residues: 1-13 <TSU>
A;Cross-references: UNIPARC:UPI000017B121
A;Experimental source: callus
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C;Species: Locusta migratoria (migratory locust)
C;Date: 21-Jul-1995 #sequence_revision 28-Jul-1995 #text_change 09-Jul-2004
C;Accession: B61620
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Pred. No. 2e+03;
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A; Cross-references: UNIPARC: UPI000017B0AD
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Search completed: January 26, 2006, 08:05:03 Job time : 9.5 secs

Run

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Result
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US-10-431-38-10
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US-10-431-638-3
US-10-431-638-3
US-10-431-638-39
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Sequence 4, Application US/11145035
Publication No. US20050287122A1
GENERAL INFORMATION:
APPLICANT: Bartlett et al.
TITLE OF INVENTION: AAV VECTORS AND METHODS
FILE REFERENCE: 28335/41335
CURRENT APPLICATION NUMBER: US/11/145,035
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 10/038,972
PRIOR PILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: US 60/260,124
PRIOR APPLICATION NUMBER: US 60/260,124
PRIOR APPLICATION NUMBER: US 60/260,124
PRIOR PILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 45
SOPTWARE: PatentIn version 3.3
SEQ ID NO 4
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OTHER INFORMATION: Synthetic peptide
US-11-145-035-4
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NUMBER
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Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2757
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind FILE REFERENCE: PF523P3
                                                      PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
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TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                               OR APPLICATION NUMBER: 60/580,347
OR FILING DATE: 2004-06-18
OR APPLICATION NUMBER: 10/29,418
OR APPLICATION NUMBER: 10/29,418
OR APPLICATION NUMBER: 60/31,469
OR FILING DATE: 2001-11-16
OR APPLICATION NUMBER: 60/340,817
OR APPLICATION NUMBER: 09/880,748
OR APPLICATION NUMBER: 09/880,748
                                                                                             APPLICATION NUMBER: 60/277,379
FILLING DATE: 2001-03-21
APPLICATION NUMBER: 60/276,248
FILING DATE: 2001-03-16
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US-10-859-643-762
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                                                    CURRENT APPLICATION NUMBER: US/11/097,864
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR FILING DATE: 2001-11-07
                                                                                                                                                                      APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 161P2F10B
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 511582006205
NUMBER OF SEQ ID NOS: 765
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 762
                                                                                                                                                                                                                                                                                                             APPLICANT: Challita-Eid, Pia M. APPLICANT: Raitano, Arthur B.
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CURRENT FILING DATE: 2004-06-02
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TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Entitled 161P2F10B Useful in Treatment and Detection
TITLE OF INVENTION: Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Challita-Eid, Pia M. APPLICANT: Raitano, Arthur B. APPLICANT: Faris, Mary
                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 10/005,480 PRIOR FILING DATE: 2001-11-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 511582006203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 16
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LLTGYPFY 12
                                                                                                                                                                                                                                                                        Hubert, Rene S.
                                                                                                                                                                                                                                                                                           Faris, Mary
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62.5%;
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Pred. No.
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15;
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US-11-097-912-762
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; TYPE: PRT
; ORGANISM: Homo Sapien
US-11-097-864-762
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US-11-097-912-762
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CURRENT APPLICATION NUMBER: US/11/097,912
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 765
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SEQ ID NO 762
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Best Local Similarity 46.2%;
Matches 6; Conservative
APPLICANT: Grey, Howard M.
APPLICANT: Epimmune Inc.
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007
CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
                                                                                                                                                                                                                                                               APPLICANT:
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TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED 161P2F10B
TITLE OF INVENTION: USEFUL IN TREATMENT AND DETECTION OF CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Challita-Eid, Pia M. APPLICANT: Raitano, Arthur B. APPLICANT: Faria, Mary APPLICANT: Hubert, Rene S.
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                                                                                                                                                                                                                                                                                                                        INFORMATION:
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                                                                                                                                                                                                                                                             Sidney, John
Southwood, Scott
                                                                                                                                                                                    Chesnut, Robert
Baker, Denise Marie
Celis, Esteban
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Morrison, Karen Jane Meyrick
                                                                                                                                                                      Kubo, Ralph
                                                                                                                                                                                                                                               Livingston,
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D. US20050265921A1
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Pred. No.
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RESULT 7
US-10-665-658-53
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PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 14490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 53, Application No. US2005
Publication No. US2005
GENERAL INFORMATION:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 53
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                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/031971
FILING DATE: 27-NCV-1996
APPLICATION NUMBER: 08/974899
FILING DATE: 20-NCV-1997
APPLICATION NUMBER: 09/420745
PILING DATE: 20-OCT-1999
PILING
                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 09/975798
FILING DATE: 28-FEB-2001
ATTORNEY/AGENT INFORMATION:
                                                                                                        NAME: Tan, Lee K.

REGISTRATION NUMBER: 39,447

REFERENCE/DOCKET NUMBER: P1014R1C1D1C1

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compartible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Windatin (Genentech)
CURRENT APPLICATION DATA.
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FILING DATE: 1993-11-29
APPLICATION NUMBER: US 08/159,339
FILING DATE: 1993-11-29
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COMPUTER READABLE FORM:
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ADDRESSEE: Genente
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APPLICATION NUMBER: US 08/073,205
FILING DATE: 1993-06-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jardieu, Paula M.
Presta, Leonard G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 DSILGGVATYGAA 16
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                                  TELEPHONE: 650/225-4462
TELEFAX: 650/952-9881
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ilarity 46.2%;
Conservative
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Pred. No. 16;
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                                                            Query Match
Best Local S
Matches 5
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Best Local :
                                                                                                                                                                                                    REGISTRATION NUMBER: 39,447
REFERENCE/DOCKET NUMBER: P1014R1C1D1C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-4462
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 57:
SEQUENCE CHARACTERISTICS:
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Presta, Leonard G.
TITLE OF INVENTION: Humanized Anti-CDlla Antibodies
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                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/031971
FILING DATE: 27-NOV-1996
APPLICATION NUMBER: 00/974899
FILING DATE: 20-NOV-1997
APPLICATION NUMBER: 09/420745
FILING DATE: 20-OCT-1999
APPLICATION NUMBER: 09/975798
FILING DATE: 28-FEB-2001
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                                                            Similarity 5; Conserv
GIYFYGTA 8
                               GVPFYGRA 13
                                                                                                                                                       LENGTH: 11 amino acids
TYPE: Amino Acid
TOPOLOGY: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/10/665,658 FILING DATE: 19-Sep-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: California
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TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                 NAME: Tan, Lee K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: 3.
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                                                            Score 28; DB
Pred. No. 16;
1; Mismatches
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Pred. No. 16;
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RESULT 9
US-10-665-658-54
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US-11-145-035-2
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APPLICANT: Bartlett et al.
TITLE OF INVENTION: AAV VECTORS AND METHODS
FILE REFERENCE: 2835/41335
CURRENT APPLICATION NUMBER: US/11/145,035
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 10/038,972
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: US 60/260,124
PRIOR FILING DATE: 2001-01-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-665-658-54
                                                                                                                                                                                         Sequence 2, Application US/11145035 Publication No. US20050287122A1 GENERAL INFORMATION:
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Publication No. US20050276801A1
GENERAL INFORMATION:
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Best Local Similarity
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REFERENCE/DOCKET NUMBER: P1014R1C1D1C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-4462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 09/975798
FILING DATE: 28-FEB-2001
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AL INFUNERAL APPLICANT: Jardieu, Paura .
APPLICANT: Dresta, Leonard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 54:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
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FILING DATE: 27-NOV-1996
APPLICATION NUMBER: 08/974899
FILING DATE: 20-NOV-1997
APPLICATION NUMBER: 09/420745
FILING DATE: 20-OCT-1999
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                                                                                                                                                                                                                                                                                                                                               GIAFYG 6
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TYPE: Amino Acid
TOPOLOGY: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/10/665,658 FILING DATE: 19-Sep-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: <Unknown>
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Pred. No. 24;
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US-11-145-035-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Synthetic Bovine Papilloma Virus Peptide US-11-145-035-6
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; OTHER INFORMATION: Synthetic Bovine Papilloma Virus Peptide
US-11-145-035-2
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Best Local Similarity
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Best Local Similarity
Matches 5; Conserv
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 2
LENGTH: 11
                                         APPLICANT: DUFFÍN, DAVID J.

APPLICANT: PIPLANI, SUNILA
APPLICANT: MERRILL, BARBARA MCEOWEN
APPLICANT: MERRILL, BARBARA MCEOWEN
APPLICANT: SCHATZ, PETER JOSEPH
TITLE OF INVENTION: COMPOUNDS HAVING AFFINITY FOR THE GRANULOCYTE-COLONY
TITLE OF INVENTION: USE
TITLE OF INVENTION: USES
FILE REFERENCE: 0300-0014
CURRENT APPLICATION NUMBER: US/10/929,988
CURRENT APPLICATION NUMBER: US/99/620,091
PRIOR APPLICATION NUMBER: US/09/620,091
PRIOR FILING DATE: 2000-07-20
NUMBER OF SEC IN NOC. 407-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Bartlett et al.
TITLE OF INVENTION: AAV VECTORS AND METHODS
FILE REPERENCE: 28335/41335
CURRENT APPLICATION NUMBER: US/11/145,035
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 10/038,972
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: US 60/260,124
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                                                                                                                                                                                                                                             APPLICANT: CWIRLA, STEVE APPLICANT: BALU, PALANI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial sequence
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ORGANISM: Artificial sequence
US/0.
UP SEQ ID NOS: 491
T: PatentIn Ver
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50.0%;
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Pred. No. 5.8e+04;
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Pred. No.
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US-11-139-435-4
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                                            OTHER INFORMATION:
                                                               FEATURE:
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APPLICANT: Fann, Ming-Ji
TITLE OF INVENTION: Cellular Proliferation Co
FILE REFERENCE: 17741-002001
CURRENT APPLICATION NUMBER: US/11/139,435
CURRENT FILING DATE: 2005-05-27
PRIOR APPLICATION NUMBER: US 60/575,611
PRIOR FILING DATE: 2004-05-27
NUMBER OF SEQ ID NOS: 36
                                                        SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 19
TYPE: PRT
                                                                                                                                                                                                                                                                                        Sequence 4, Application US/11139435 Publication No. US20050287664A1 GENERAL INFORMATION:
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Publication No. US20050287122A1
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CURRENT APPLICATION NUMBER: US/11/145,035
CURRENT FILING DATE: 2005-06-03
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PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: US 60/260,124
PRIOR FILING DATE: 2001-01-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 11
TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
FOTHER INFORMATION: Synthetic Bovine Papilloma Virus Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bartlett et al.
TITLE OF INVENTION: AAV VECTORS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
                                    ORGANISM: Artificial Sequence
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conditioned
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 26; DB
Pred. No. 36;
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Pred. No. 32;
medium (CM)
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US-10-665-658-56
; Sequence 56, Application US/10665658
; Publication No. US20050276801A1
; GENERAL INFORMATION:
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                                                                                  RESULT 16
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Publication No. US2005
GENERAL INFORMATION:
APPLICANT: Jard
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Best Local
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Best Local
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REFERENCE/DOCKET NUMBER: P1014R1C1D1C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-4462
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/031971
FILING DATE: 27-NOV-1996
APPLICATION NUMBER: 08/974899
FILING DATE: 20-NOV-1997
APPLICATION NUMBER: 09/420745
FILING DATE: 20-OTT-1999
APPLICATION NUMBER: 09/975798
FILING DATE: 28-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/665,658
FILING DATE: 19-Sep-2003
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
APPLICANT: Jardieu, Paula M.
                                                                                                                                                                                                                                                                                      SEQUENCE DESCRIPTION: SEQ ID NO: 51:
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                                                                                                                                                                                                    Similarity
4; Conserv
                                                                                                                                      GIYFYG
                                                                                                                                                                                                                                                                                                      LENGTH: 11 amino acids
TYPE: Amino Acid
TOPOLOGY: <Unknown>
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CITY: South San Francisco
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Presta, Leonard G.
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42.9%;
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RESULT 17
US-10-665-658-58
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 27-NOV-1996
APPLICATION NUMBER: 08/974899
FILING DATE: 20-NOV-1997
APPLICATION NUMBER: 09/420745
FILING DATE: 20-OCT-1999
APPLICATION NUMBER: 09/975798
FILING DATE: 28-FEB-2001
ATTORNEY/AGENT INFORMATION:
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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APPLICATION NUMBER: US/10/665,658
FILING DATE: 19-Sep-2003
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
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TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 39,447
REFERENCE/DOCKET NUMBER: P1014R1C1D1C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-4462
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ADDRESSEE: Genente
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                                                                                                                                                                                                                                                                                                                        APPLICANT: Jardieu, Paula M.
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                                                                                                                                                      CITY: South San F:
STATE: California
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TOPOLOGY: <Unknown>
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STREET: 1 DNA Way
                                                                                                                                     COUNTRY: USA
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                                                                                                                                                                              Francisco
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Pred. No. 54;
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RESULT 18
US-10-665-658-12
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Publication No. US20050276801A1
GENERAL INFORMATION:
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Best Local
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INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: Amino Acid
             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinBatin (Genentech)
CURRENT APPLICATION NOWBER: US/10/665,658
FILING DATE: 19-Sep-2003
CLASSIFICATION NUWBER: 05/031971
APPLICATION NUWBER: 60/031971
APPLICATION NUWBER: 60/031971
FILING DATE: 27-NOV-1996
APPLICATION NUWBER: 08/974899
FILING DATE: 20-NOV-1997
APPLICATION NUWBER: 09/97499
FILING DATE: 20-NOV-1997
APPLICATION NUWBER: 09/975798
FILING DATE: 20-NOV-1997
APPLICATION NUWBER: 09/975798
FILING DATE: 20-NOV-1997
APPLICATION NUWBER: 09/975798
FILING DATE: 20-FEB-2001
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REGISTRATION NUMBER: 39,447
REFERENCE/DOCKET NUMBER: P1014R1C1D1C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-4462
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  ATTORNEY/AGENT
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CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Presta, Leonard G.
TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Jardieu, Paula M.
Presta, Leonard C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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FILING DATE: 27-NOV-1996
APPLICATION NUMBER: 08/974899
FILING DATE: 20-NOV-1997
APPLICATION NUMBER: 09/420745
FILING DATE: 20-OCT-1999
APPLICATION NUMBER: 09/975798
FILING DATE: 28-FEB-2001
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STATE: California
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INFORMATION:
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                                                                                                                                          ORGANISM: Homo sapiens
US-11-054-515-2893
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                                                                                                                                                                                                            SEQ ID NO 2893
                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Publication No. US20
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 30.1%;
Best Local Similarity 66.7%;
Matches 4; Conservative
                                                                                                       Query Match
                                                                                        Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/543,296 PRIOR FILING DATE: 2004-02-11
                                                                                                                                                                                                                             Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 3247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                             LENGTH: 16
TYPE: PRT
                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-10-17
                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR
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INFORMATION FOR SEQ ID NO: 12:
                                                                                      Local
                                                                                                                                                                                                                                                                                                   FILING DATE: 2001-03-21
APPLICATION NUMBER: 60/
FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 09/880,748
FILING DATE: 2001-06-15
APPLICATION NUMBER: 60/293,499
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/331,469 FILING DATE: 2001-11-16
                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/240,816
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/277,379
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/340,817 FILING DATE: 2001-12-19
                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 39,447
REFERENCE/DOCKET NUMBER: P1014R1C1D1C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-4462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE DESCRIPTION: SEQ ID NO:
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                                   3 LVMGVPFYG 11
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                                                                                    Similarity
ILTĠŸYYYĠ 13
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TOPOLOGY: Linear
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                                                                      Conservative
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                                                                                    Score 25;
Pred. No.

    Mismatches

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Pred. No. 60;
                                                                      Mismatches
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; Sequence 9, Application US/10431638 ; Publication No. US20060003939A1

RESULT 20 US-10-431-638-9

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APPLICANT: The Rockefeller Institute
APPLICANT: Steller, Hermann
ITITLE OF INVENTION: PEPTIDES AND METHODS FOR
FILE REFERENCE: P-5004-US
CURRENT APPLICATION NUMBER: US/10/431,638
CURRENT FILING DATE: 2003-05-08
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 7
TYPE: PRT
ORGANISM: Drosophila melanogaster
                                                                                                                                                                                                                                 Sequence 597, Application US/10859
Publication No. US20060002993A1
GENERAL INFORMATION:
APPLICANT: Challita-Eid, Pia M.
APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
APPLICANT: Hubert, Rene S.
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US-10-859-643-597
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US-10-431-638-10
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Publication No. US20060003939A1
GENERAL INFORMATION:
APPLICANT: The Rockefeller Institute
APPLICANT: Steller, Hermann
TITLE OF INVENTION: PEPTIDES AND METHODS FOR CELL DEATH REGULATION
FILE REFERENCE: P-5004-US
CURRENT APPLICATION NUMBER: US/10/431,638
CURRENT FILING DATE: 2003-05-08
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.3
SEQ ID NO 10
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            APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Entitled 161P2F10B Useful in Treatment of Invention: Cancer
FILE REFERENCE: 511582006203
CURRENT APPLICATION NUMBER: US/10/859,643
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR PILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 765
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TYPE: PRT
ORGANISM: Drosophila melanogaster
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FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                              Morrison, Karen Jane Meyrick
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Faris, Mary
Hubert, Rene S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28.9%; Score 24; DB 6; Lo
100.0%; Pred. No. 5.8e+04;
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APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Entitled 161P2F10B Useful in Treatment at
TITLE OF INVENTION: Cancer
FILE REFERENCE: $11582006203
CURRENT APPLICATION UNMBER: US/10/859,643
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR FILING DATE: 2001-11-07
NUMBER OF 58Q ID NOS: 765
NUMBER OF SRG ID NOS: 765
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Challita-Eid, Pia M.
APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
APPLICANT: Hubert, Rene S.
APPLICANT: Morrison, Karen Jane
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; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-859-643-597
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US-10-859-643-685
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Best Local Similarity
                                                                                                                                                         APPLICANT: Hubert, Rene S.

APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 161P2F10B
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 511582006205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Challita-Eid, Pia M. APPLICANT: Raitano, Arthur B. APPLICANT: Faris, Mary APPLICANT: Hubert, Rene S.
             CURRENT APPLICATION NUMBER: US/11/097,864
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 765
SOFTWARE: FastSEQ for Windows Version 4.0
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ORGANISM: Homo Sapien
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Pred. No. 5.8e+04;
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Pred. No. 5.8e+04;
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APPLICANT: Paris, Mary
APPLICANT: Hubert, Rene S.
APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 161P2F10B
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF CANCER
FILLE REPERENCE: 51158206205
CURRENT APPLICATION NUMBER: US/11/097,864
CURRENT APPLICATION NUMBER: US/10/062,109
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 765
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 685
LENGTH: 9
TYPE: PRT
ORGANISM: Homo Sapien
US-11-097-864-685
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; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo Sapien
US-11-097-864-597
Search completed: January 26, 2006, 08:39:07 Job time : 4.22414 secs
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US-11-097-864-685
                                                                                                                                                                                       Query Match 28.9%; Score 24; DB 7; Length 9; Best Local Similarity 44.4%; Pred. No. 5.8e+04; Matches 4; Conservative 3; Mismatches 2; Indels
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APPLICANT: Raiteno, Arthur B.
APPLICANT: Faris, Mary
APPLICANT: Hubert, Rene S.
APPLICANT: Morrison, Karen January
                                                                                              7 VPFY 10
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Q9989 HUMAN
Q4YM11 PLABE
Q74UV1 PLABE
Q74UV1 PLABE
Q74UV1 PLABE
Q74UV1 PLACH
Q95830 PINMO
TY13 BÖMVA
Q15632 HUMAN
Q34K15 PLACH
Q64D23 MOUSE
Q76103 Q8YSA
Q4XLTS PLACH
Q64D23 MOUSE
Q771013 ORYSA
Q4XLTS PLACH
Q64D23 MOUSE
Q77101 NEUCR
Q8AXQ7 XENLA
Q98XQ0 HUMAN
Q7TSR9 MOUSE
Q7M2G7 VICFA
Q64X1K5 MOUSE
Q7M2G7 VICFA
Q8AXGT HESAL
TYLP3 SPIOL
Q9TWN3 THESE
P81896 SOLTU
TL19 ARATH
Q7RZ51 NEUCR
Q9RF21 MYCMY
Q80817 9DELA
Q51S44 9ARAE
Q6JU97 NEIGO
Q50019 MYCLE
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  Q99898 homo
Q4yw11 plasm
Q7auj5 acine
P20016 thunn
Q9880 pinus
P84215 bombi
Q15632 homo
Q98818 brass
Q4x1t5 plasm
Q6123 mus m
Q7m1u3 oryza
Q782j9 neuro
Q8axq7 xenop
Q9bxq0 homo
Q7tsr9 mus m
Q7m1u3 rhodo
Q8x3k5 mus m
P82799 spina
Q61dn4 rhodo
Q8x3k5 mus m
P83149 rhodo
Q8x3k5 mus m
P83149 spina
Q9twn3 theil
P8199 spina
Q9twn3 theil
P8199 spina
Q9twn3 theil
P8199 spina
Q9twn3 theil
P8199 spina
Q9twn3 theil
Q9tz1 meuro
Q9rfz1 mycop
Q9rf21 mycop
Q5041 neiss
Q5141 neiss
 mycoplasma
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                                          PSAL_SYNVU
Q9UMFIO_9EURY
Q8NEDS HUMAN
Q96AH6_HUMAN
Q96AH6_HUMAN
Q96AH6_PARA
Q88250_PARA
Q89254_PARA
ANGTI BOTJA
TKUI ÜREUN
Q7M271_SHEEEN
Q7M271_SHEED
H2AX ONCMY
Q7LZ51_PHYBI
TKNI_SCHGR
Q7M271_BOMMO
Q9UMCI_9EURY
Q7M144_RAPAH
Q7M285_SOYEN
Q35188_TOBAC
Q9URCI_SALTI
Q9PR29_MICUN
TRYP_FELCA
Q4M373_9LECA
Q4M373_9LECA
Q4M373_9LECA
Q4M373_9LECA
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SOVO_SEPOF
QAY/B4 PLACH
QBH229_9VIRU
QBQUVI_9WURI
QQQVIP
QAY/B4 PLAYO
QRYRB3 BORBU
QRYRB9 BORBU
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QRYRB9 BORBU
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QRYRB1 PLAWA
AZOUI
FLAW AZOUI
CAMPA
QRYRB9 BORBU
QRYB9 BORBU
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Q457547
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arabidopsis
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RESULT 1
Q99898 HUMAN E
ID Q59898 HUMAN E
AC Q99898;
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DT 01-MAY-199
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Q4YW11;
13-SEP-2005 (7
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata
Eukaryota; Eutheria; Euarchontoglires;
                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.
Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Hall N., Karras M., Raine J.D., Carlton J.M., Christophides G.K.,
Berriman M., Florens L., Janssen C.S., Pain A., Churcher C.,
James K., Rutherford K., Harris B., Harris D., Churcher C.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.C.,
Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.;
"A comprehensive survey of the Plasmodium life cycle by genomic,
transcriptomic, and proteomic analyses.";
Science 307:82-86(2005).
-i- CAUTION: The sequence shown here is derived from an
EMBL/Genbank/DDBJ whole genome shotgun (WGS) entry which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
Hypothetical protein (Fragment).
ORFNames=PB105279,00.0;
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Eukaryota; Alveolata;
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01-MAY-1997 (TrEMBLrel. 03, Last
01-MAR-2001 (TrEMBLrel. 16, Last
Mutant Sos isoform 1 (Fragment).
                                                                                                                                                   SEQUENCE
                                                                                                                                                                                      Hypothetical
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                                                                                                                                                                                                                                                                preliminary data.;; CAAI01002172; (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TaxID=5821;
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16 AA;
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Pred. No. 1.3e+03;
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Matches 4
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01-FEB-1991 (Rel. 17, Last sequence t
10-MAY-2005 (Rel. 47, Last annotation
NAD-specific glutamate dehydrogenase
(Fragment).
                                                                                                                                                                                                                                       DHE2_THUTH P20016;
MEDLINE=76253947; PubMed=182270; DOI=10.1016/0005-2744 (76) 90156-X; Veronese F.M., Bevilacqua R., Boccu E., Brown D.M.; "Purification, characteristics and sequence of a peptide containing essential lysine residue."; Biochim. Biophys. Acta 445:1-13(1976).
                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Percifor
Scombridae; Thunnus.
NCBI_TaxID=8237;
                                                                                                                                                                                                                                                               HITUHI
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (MAY-2002) to the EMBL; AJ486856; CAD31077.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transposons in the natural environment: study of adjacent DNA sequences."; microbiology 150:979-992(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmid pKLH207.
Bacteria; Proteobacteria; Gam
Moraxellaceae; Acinetobacter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                               Thunnus thynnus (Bluefin tuna)
                                                                                                                                                                                                                                                                                                                                                                                                                              Plasmid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nikiforov V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=15073307; DOI=10.1099/mic.0.26844-0; Kholodii G., Mindlin S., Gorlenko Z., Petrova
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN-BW3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=tnp1006;
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                                                                      TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kholodii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Translocation of transposition-deficient [Tn(d)PKLH2-like]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Acinetobacter sp. BW3
                                                                                  PROTEIN SEQUENCE.
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4; Conservative
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2 AA;
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80.0%;
 L-glutamate + H(2)O + NAD(+) = 2-oxoglutarate
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-; Genomic_DNA.
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Pred. No. 2.
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                                                                                                                              Percomorpha; Perciformes; Scombroidei;
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(EC 1.4.1.2)
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RESULT 6
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01-FEB-2005 (Rel. 4
01-FEB-2005 (Rel. 4
13-SEP-2005 (Rel. 4
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Plant Cell
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ribulose BIPHOSPHATE carboxylase small subunit (Fragment). Pinus monticola (Western white pine). Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheo Spermatophyta; Coniferopeida; Coniferales; Pinaceae; Pinus; Ini TaxID-3345;
                                        Tryptophyllin 13.
Bombina variegata (Yellow-bellied toad).
Bukaryota; Metazoa; Chordata; Craniata;
Eukaryota, Batrachia; Anura; Archeobatrac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=96213005; Pu
Ekramoddoullah A.K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13, 01-JUN-2000 (TrEMBLrel. 14,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PROTEIN SEQUENCE.
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Q9S8A0;
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PROTEIN SEQUENCE,
                                  NCBI_TaxID=8348;
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InterPro; IPR060697; GLFV_dehydrog_N.
Pfam; PF02812; ELFV_dehydrog_N.
PROSITE; PS00074; GLFV_DEHYDROGENASE; PARTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIR; A12729; A12729.
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FUNCTION,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=8665095;
K., Taylor D.W.;
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1803 MW; 27D4934C881717DE
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                                                            a; Craniata; Vertebrata; Euteleostomi;
Archeobatrachia; Bombinatoridae; Bombina.
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SUBCELLULAR LOCATION,
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Pred. No.
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"Skin secremian"
                                                                                                                                                                                                                                                           MEDLINE=90222189; PubMed=2326274;
Zutter M., Hockett R.D., Roberts C.W.,
Morton C.C., Deaven L.L., Crist W.M., C
"The t (10;14) (q24;q11) of T-cell acute
juxtaposes the delta T-cell receptor wi
                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (Human).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Q15632;
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SEQUENCE 14 AA; 1651 MW;
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                                                              Local
                                                                                   Match
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GO:0005179; F:hormone activity; TAS.
GO:0006952; P:defense response; TAS.
GO:006956; P:regulation of insulin secretion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98
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Pred. No. 5.6e
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RESULT 10
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
Chalcone synthase (Fragment).
Arabis pauciflora.
Eukaryota · V'-'''.
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Q9AUA1;
01-JUN-2001 (;
01-JUN-2001 (;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROTEIN SEQUENCE.

MEDLINE-95218603; PubMed=7766044; DOI=10.1007/BF00202596; Falk A., Taipalensuu J., Ek B., Lenman M., Rask L.; Planta 195:387-395(1995).

InterPro; IPR001229; Jacalin_lectin.
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Q9S8H8;
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Brassica napus (Rape).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;

eurosids II; Brassicales; Brassicaceae; Brassica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UN-2002 (TrEMBLrel. 21, Last annotation update)
Myrosinase-binding protein 50 (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=3708;
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SEQUENCE 16 AA; 1763 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Koch M., Kroymann J., Haubol Submitted (MAR-2000) to the EMBL; AF248998; AAK31933.1; NON TER 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=21441848; PubMed=11557794;
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NCE 20 AA; 2101 MW;
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EMBL/GenBank/DDBJ databases
-; Genomic_DNA.
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eudicots; rosids;
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Q6LD23_MOUSE
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Best Local
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MEDLINB=95047447; PubMed=7958966; DOI=10.1016/0378-1119(94)90710-2;
Lutfalla G., Uze G.;
"Structure of the murine interferon alpha/beta receptor-encoding gene:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence up
05-JUL-2004 (TrEMBLrel. 27, Last annotation
Interferon alpha/beta receptor (Fragment).
Name=Ifnarl; Synonyms=IFNAR;
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Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Hall N., Karras M., Raine J.D., Carlton J.M., Christophides G.
Berriman M., Florens L., Janssen C.S., Pain A., Christophides G.
James K., Rutherford K., Harris B., Harris D., Churcher C.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos
Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.;
"A comprehensive survey of the Plasmodium life cycle by genomic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence up
13-SEP-2005 (TrEMBLrel. 31, Last annotation
13-SEP-2005 (TrEMBLrel. 31, Last annotation
Hypothetical protein (Fragment).
ORFNames=PC109111.00.0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
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Eukaryota; Alveolata; Apicomplexa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                            EMBL; U06237; AAA65003.1; ~; MGI; MGI:107658; Ifnar1.
                                                                                                                                                                                                                                                                                                                                            Gene 148:343-346(1994).
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Galagan J.E., Calvo S.E., Borkovich K.A., Selker E.U., Read N.D.,

Jaffe D., Fitzhugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,

Jaffe D., Fitzhugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,

Rifins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,

A Lakins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,

A Cui D., Ianakiev P., Pedersen D., Nelson M., Washburne M.,

Selitrennikoff C.P., Kinsey J.A., Braun E.L., Zelter A., Schulte U.,

Kothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,

A Kothe G.O., Jedd G., Newes W., Staben C., Rudd S., Frishman D.,

Roy A., Foley K., Naylor J., Thomann N., Barrett R., Gnerre S.,

A Roy A., Foley K., Naylor J., Thomann N., Barrett R., Greenberg D.,

A Roy A., Kamal M., Kamvysselis M., Mauceli E., Bielke C., Rudd S., Frishman D.,

Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S.,

A Cogoni C., Macino G., Catcheside D., Li W., Pratt R.J., Osmani S.A.,

DeSouza C.C., Glass L., Orbach M.J., Berglund J., Voelker R.,

Varden O., Plamann M., Seiler S., Dunlap J., Radford A., Aramayo R.,

Paulsen I., Sachs M.S., Lander E.S., Nusbaum C., Birren B.,

"The Genome Sequence of the Filamentous Fungus Neurospora crassa.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NON TER
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Q752J9;
                                             Nature 0:0-0(2003).
-|- CAUTION: The sequence sh
EMBL/GenBank/DDBJ whole
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01-MAR-2004
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Name=NCU04977.1;
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"A rice protein library; a data-file of
two-dimensional electrophoresis.";
Theor. Appl. Genet. 86:935-942(1993).
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     EAA29651.1;
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Genomic_DNA
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SEQUENCE
               Citron B.A., Santa Cruz K.S., Davies P.J.
"Intron-exon swapping of tranglutaminase
aggregation in Alzheimer's disease.";
J. Biol. Chem. 0:0-0(2001).
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Endo Y., Nonaka M., Saiga H.,
Matsushita M., Fujita T.;
"Origin of mannose-binding le
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01-MAR-2003 (TrEMBLrel. 23,
01-OCT-2003 (TrEMBLrel. 25,
                                                                      TISSUE=Brain;
Citron B.A.,
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Q8AXQ7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    igin of mannose-binding lectin-associated MASP-3 involved in the lectin complement invertebrate, amphioxus "; Immunol. 170:4701-4707(2003).
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 AF311286;
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| GVPRFSRS
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Eutheria;
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Euarchontoglires; Primates; Catarrhini;
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Pred. No. 7.7e+03
2; Mismatches
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tivity; TEA.
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1. No. 8.3e+03;
1. No. 8.3e+03;
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RESULT
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Best Loc
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Best Local
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GO; GO:0005524; F
GO; GO:0005525; F
GO; GO:0003924; F
GO; GO:0006916; P
GO; GO:0019933; P
GO; GO:0019221; P
GO; GO:0007186; P
GO; GO:0018276; P
GO; GO:0012501; P
ACY1transferase; T
                                                                                                                        Q7M2G7_VICFA PRELIMINARY;
Q7M2G7;
Q1MAR-2004 (TrEMBLrel. 26,
Q1-MAR-2004 (TrEMBLrel. 26,
Q1-MAR-2004 (TrEMBLrel. 26,
Q1-MAR-2004 (TrEMBLrel. 26,
NADH2 dehydrogenase (EC 1.0,
Vicia faba (Broad bean).
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Q7TSR9_M
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Mammalia; Eutheria; Euarchontoglires;
Muridae; Murinae; Mus
                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tre
Spermatophyta; Magnoliophyta; eudicotyledons; core eudic
eurosids I; Fabales; Fabaceae; Papilionoideae; Vicieae;
NCBI_TaxID=3906;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=C57BL/6;
Fujino T., Lee W.-C.A.,
"Regulation of cpg15 by
Plasticity.";
                                                                                                        Mitochondrion.
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01-OCT-2003
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VICFA
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3L; AY150584; AAN84528.1; -; Genomic_DNA.
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GO:0005524; F:ATP binding; TAS.
GO:0005525; F:CTP binding; TAS.
GO:0005525; F:CTP binding; TAS.
GO:0003924; F:GTPase activity; TAS.
GO:0003810; F:protein-glutamine gamma-glutamyltransferase.
GO:0006916; F:anti-appytosis; TAS.
GO:0019933; P:CAMP-mediated signaling; TAS.
GO:0019931; P:cytokine and chemokine mediated signaling p.
GO:0007166; F:Gpyrotein coupled receptor protein signalin.
GO:0018276; P:isopeptide cross-linking via N6-glycyl-L-ly.
GO:0012501; P:programmed cell death; TAS.
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Signaling
                                                              Streptophyta; Embryophyta; Tracheophyta; yta; eudicotyledons; core eudicots; rosids;
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Pred. No. 9e+03;
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1.1e+04;
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RESULT 19
QBKXK5 MOUSE PRELIMINARY;
ID QBKXK5;
AC QBKXK5;
DT 01-OCT-2002 (TrEMBLrel. 22,
DT 0
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05-JUL-2004 (TrEMBLrel. 27, Last sequence up
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05-JUL-2004 (TremBLrel. 27, Last annotation
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NON TER
SEQUENCE
                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Euarchontoglires; Glires; Rodent Muridae; Murinae; Mus.
                                                                                                                                                                                                                      01-OCT-2002 (TrEMBLrel. 22, 01-OCT-2002 (TrEMBLrel. 22, 01-OCT-2002 (TrEMBLrel. 22, Arginase I (Fragment).
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=88313675; PubMed=3410321; DOI=10.1016/0378-1119(88)90463-5; Pollock D., Bauer C.E., Scolnik P.A.; "Transcription of the Rhodobacter capsulatus nifhDK operon is modulated by the nitrogen source: Construction of plasmid expression vectors based on the nifhDK promoter."; Gene 65:269-275(1988).

Gene 65:269-275(1988).

Gene 65:269-275(1988).

Gene 65:269-275(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R.capsulata nifHDK operon, (Fragment).
Rhodobacter capsulatus (Rhodopseudomonas capsulata).
Bacteria; Proteobacteria; Alphaproteobacteria; Rhodobacterales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IT 18
14_RHOCA
Q6LDN4_RHOCA PRELIMINARY;
Q6LDN4;
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"Purification and preliminary characterization of mitochondrial complex I (NADH:ubiquinone reductase) from broad bean (Vicia fa
  NUCLEOTIDE SEQUENCE.
STRAIN=129/SvJ;
PubMed=15922518; DOI=10.1016/j.gene.2005.04.004;
Gray M.J., Poljakovic M., Kepka-Lenhart D., Morr:
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PIR; PQ0790; PQ0790.
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9_SPIOL
TU19_SPIOL
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STRAIN-NCIMB 1903T;
Onji M., Hirabayashi J., Suzuki S.;
Onji M., Hirabayashi J., Suzuki S.;
Onji M., Hirabayashi J., Suzuki S.;
Characterization of major outer membrane proteases.";
Microbes Environ. 0:0-0(2002);
Microbes Environ. 0:0-0(2002);
Microbes Environ. (By similarity).

-1- SUBUNIT: Homotrimer (By similarity).
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                                                                                       P82799;
16-OCT-2001
Spermatophyta; Magnoliophyta; eudicotyle Caryophyllales; Amaranthaceae; Spinacia. NCBI TaxID=3562;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Induction of arginase I transcription by IL-4 requires a DNA response element for STAT6 and C/EBPbeta."; Gene 353:98-106(2005).

EMBL; AY074884; AAL79623.1; -; Genomic_DNA.

NON TER

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                                                                  16-OCT-2001 (Rel.
16-OCT-2001 (Rel.
05-JUL-2004 (Rel.
                                                                                                                                                                                                                                                   Direct protein sequencing; Ion transport; Porin; Transmembrane; Transport.

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28-PEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation updat
Outer membrane protein 40Va (Omp40Va) (Frag
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P83149;
                                      Spinacia oleracea (Spinach)
Eukaryota, Viridiplantae, S
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                                                            Thylakoid lumenal
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Vibrionaceae; Vibrio.
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40, Last sequence update)
44, Last annotation update)
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yta; eudicotyledons;
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Pred. No. 1.2e
1; Mismatches
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RESULT 23
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Q9TWN3_THESE
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01-JUN-2000
01-JUN-2003
                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embry
Spermatophyta; Magnoliophyta; eudicotyledons;
lamiids; Sólanales; Solanaceae; Solanum.
                                                                                                                                Dihydrolipoamide succinyltransferase dehydrogenase complex (EC 2.3.1.61) (Solanum tuberosum (Potato)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baek B.K., Kim B.S., Rhim B.M., Lee H.I., Park Y.H., Kakoma I.; "Immunogenicity and protective efficacy of solubilized merozoite-enriched Theileria sergenti immunogens. III. Characterization of
 MEDLINE=99441232;
Millar A.H., Hill
                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                              immunodominant peptides.";
Submitted (MAR-1995) to the EMBL/GenBank/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UUN-2000 (TrEMBLrel. 14, Last annotation updat
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Q9TWN3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kieselbach T., Pettersson U., Bystedt M., Schroeder W.P.;
Submitted (JUL-2000) to Swiss-Prot.
-!- SUBCELLULAR LOCATION: Chloroplast; within the thylakoid
                                              PROTEIN SEQUENCE
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                                                                                                                                                                          (TrEMBLrel. 14, Created)
(TrEMBLrel. 14, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
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; TISSUE=TUBER;
PubMed=10510296;
S.A., Leaver C.J
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Pred.
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Pred. No. 1.2e+04
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                DOI=10.1042/0264-6021:3430327
                                                                                                                                                component of 2-oxoglutarate
(E2) (OGDC-E2) (Fragment).
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dons; core e
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mitochondrial 2-oxoglutarate dehydrogenase complex:

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RESULT 24
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SEQUENCE
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16-OCT-2001 (Rel. 40, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Thylakoid lumenal 19 kDa protein (P19) (Fragment).
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicotyledons;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           purification and characterization in potato.";
Biochem. J. 343:327-334(1999).
-!- FUNCTION: THE 2-OXOGLUTARATE DEHYDROGENASE
                                                                                                                                                           Chloroplast;
                                                                                                                                                                                                                          use
                                                                                                                                                                                                                                           This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                        STRAIN=cv. Columbia;
Kieselbach T., Bystedt M., Schroeder W.P.;
Submitted (JUL-2000) to Swiss-Prot.
-!- SUBCELLULAR LOCATION: Chloroplast; within the thylakoid lumen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ARATH
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                                                                                                                                                                                                                                                                                                                                                                                                               PROTEIN SEQUENCE
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SUBCELLULAR LOCATION: MITOCHONDRIAL.

CAUTION: X IN THE SEQUENCE MIGHT BE CYS RESIDUE.

SIMILARITY: TO OTHER SPECIES E2 COMPONENT.

GO:0005739; C:mitochondrion; IEA.

GO:0008415; F:acyltransferase activity; IEA.

GO:0004149; F:dihydrolipoyllysine-residue succinyltransfe.

GO:0006470; F:transferase activity; IEA.

GO:0006096; P:glycolysis; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FUNCTION: THE 2-OXOGLUTARATE DEHYDROGENASE COMPLEX CATALYZES THE OVERALL CONVERSION OF 2-OXOGLUTARATE TO SUCCINYL-COA & CO(2). IT CONTAINS MULTIPLE COPIES OF 3 ENZYMATIC COMPONENTS: 2-OXOGLUTARATE DEHYDROGENASE (E1), DIHYDROLIPOAMIDE SUCCINYLTRANSFERASE (E2) AND LIPOAMIDE DEHYDROGENASE (E3).

CATALYTIC ACTIVITY: SUCCINYL-COA + DIHYDROLIPOAMIDE = COA + S-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SUBUNIT:
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FYGRAXS 15
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21
21 AA; 2
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                                        Conservative
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21
                                                             27.7%;
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Pred. No. 1.2e-
2; Mismatches
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Pred. No.
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1.2e+04;
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Search completed: January 26, Job time: 53.5603 secs

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RESULT 25
QTRZZ1 NEUCR PRELIMINARY;
ID QTRZZ1,
AC Q7ZZ1,
AC SCACATION COLOR
AC SCACATION COLOR
AC CSCACATION COLOR
AC CSCACATI
                                                                                                                                                                                                                                                                                                                                                                           A Galagan J.E. Calvo S.E., Borkovich K.A., Selker E.U., Read N.D., A Jaffe D., Fitzhugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B., A Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M., RA Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M., RA Gui D., Ianakiev P., Pedersen D., Nelson M., Washburne M., RA Selitrennikoff C.P., Kinsey J.A., Braun E.L., Zelter A., Schulte U., RA Koche G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D., RA Koy A., Foley K., Naylor J., Thomann N., Barrett R., Gnerre S., RA Koy A., Foley K., Naylor J., Thomann N., Barrett R., Gnerre S., RA Kamal M., Kamvysselis M., Mauceli E., Bieke C., Rudd S., Frishman D., Kamvysselis M., Mauceli E., Bieke C., Rudd S., Frishman D., RA Krystofova S., Ramussen C., Metzenberg R.L., Perkins D.D., Kroken S., Cogoni C., Macino G., Catcheside D., Li W., Pratt R.J., Osmani S.A., Pasouza C.C., Glass L., Orbach M.J., Berglund J., Voelker R., Yarden O., Plamann M., Seiler S., Dunlap J., Radford A., Aramayo R., RA Varden O., Plamann M., Seiler S., Nusbaum C., Birren B., "The Genome Sequence of the Filamentous Fungus Neurospora crassa."; RL Nature 0:0-0(2003).

"CC Cambi Genbank/DDBJ whole genome shotgun (WGS) entry which is EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
                                                                                                                                                                            Query Match
Best Local
                                                                                                                                             Matches
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Eukaryota, Fungi; Ascomycota, Pezizomycotina, Sordariomyc
Sordariomycetidae; Sordariales, Sordariaceae, Neurospora.
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                                                                                                                                                                            Similarity
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MGTKVFGR 8
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Pred. No.
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E1CE7FEAD658B6D5 (
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Purcell S., Rehman -
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1: geneseqq1980
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Gapop 10.0 ,
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109
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AAU96316

AAW18898

ABB338020

AAW11434

ABB23254

AAW71153

AAW58647

ABG452866

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ABG45286
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Abg5265
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Adc27861
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Aaw380771
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Aam31434
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Aam71153
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Aau96316
Aam18898
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Aau96324 Der HN
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AAW85207
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ABU741274
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AAB71339
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Aay45830 Immunogen
Aby14830 Immunogen
Aby14831 Immunogen
Aby15174 Immunogen
Aby15174 Immunogen
Abu170333 Human imm
Abu69779 Human imm
Abu69779 Human imm
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Abu6978 HIV-1 cro
Add96334 HIV-1 cro
Add96366 HIV-1 cro
Add9637 HIV-1 cro
Add9637 HIV-1 cro
Add9637 HIV-1 cro
Add9637 HUV-1 cro
Add9638 HIV-1 cro
Add9639 HIV-1 cro
Add9639 HIV-1 pol
Adt41877 NSARS vir
Ads80692 SARS vir
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Ads80692 FAIV DR Su
Adt11807 HSARS Vir
Ads80692 Cotton fi
Adv22436 HIV-1 Pol
Adv2436 HIV-1 Pol
Adv2436 HIV-1 Pol
Adv2436 HIV-1 Pol
Adv2436 HIV-1 HLA
Adc89725 Cotton fi
Abu89725 Cotton fi
Abu89737 Insulin/i
Ada03420 Insulin r
Add167324 IGF-1R/IR
Add04633 Insulin r
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Aap50408
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RESULT 1
AAY52520
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13-MAY-1998;
02-SEP-1998;
                                                                                                                                                                                   Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequences AAY52510-Y52522 represent proteolytic fragments of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 70; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel high molecular weight Dermatophagoides nucleic acid polypeptides used to modify an animals' hypersensitivy to mite allergens.
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98US-0085295P.
98US-0098909P.
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                                                                                                              100.0%; Score 109; DB 3
100.0%; Pred. No. 4e-11;
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Aaw88603 Secreted
Adt92805 Consensus
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RESULT 2
ARM96324
ID MAN96324
ID MAN96324
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ID ARM96324
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PM 11-1
PM MCC:
XX H-1
PM MCC:
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RESULT 3
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AC AAY5
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DE Hous
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an isolated mite allergenic protein of CC Dermatophagoides, designated Der HMW-map protein, and its related nucleic CC acid. The Der HMW-map protein is useful for eliciting an immune response CC against Der HMW-map protein. The protein or a reagent comprising a nongroup relation is useful for identifying an animal (e.g., dog, CC cat) susceptible to or having an allergic response to a mite. A CC therapeutic composition is useful for desensitising a host animal to an CC detection of anti-Der HMW-map antibodies in animal fluids, and inhibition CC disease. Antibodies that bind to Der HMW-map are useful for inhibiting confirmunoglobulin (Ig)E or Der HMW-map protein activity associated with a CC disease. Antibodies that bind to Der HMW-map are useful for inhibiting CC binding of proteins to IgE, to prevent immunocomplex formation, thus creducing hypersensitivity responses to mite allergens, and as vaccines against mite allergen hypersensitivity. Sequences AAU96314-AAU96342 cc represent Der HMW-map polypeptides of the invention
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                                                                                                                            AAY52512;
                                                                                                                                                                                      AAY52512 standard;
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Der HMW-map protein, useful as a vaccine for treating mite allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-351888/38
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House dust mite allergen protein (map) A/B fragment map(2).

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RESULT 4
AAU96316
ID AAU9
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AC AAU5
XX
DT 15--
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KW Der
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KW mite
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Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAY52510-Y52522 represent proteolytic fragments of Dermatophagoides farinae high molecular weight mite allergen -map) composition. The HMW-map tomposition was isolated from the the thing of the things o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mite allergen protein; map; high molecular weight; HMW-map; allergy; house dust mite; IgE; immunoglobulin E; allergen; mapA; mapB; hypersensitivity reaction; therapy; treatment; diagnosis; human; feline; canine; veterinary; antibody; vaccine; immunisation.
Dermatophagoides farinae
                                                        Der HMW-map; American hou mite allergenic protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     allergens from a mixture of proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel high molecular weight Dermatophagoides nucleic acid polypeptides used to modify an animals' hypersensitivy to mite allergens.
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                                                                                                                       Der HMW-map polypeptide #3
                                                                                                                                                                                                                                                   AAU96316 standard; peptide; 20
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13-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dermatophagoides farinae.
                                                                                                                                                                   15-JUL-2002
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                                         immunocomplex formation.
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19; Conservative
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                                                                                  American house
                                                                                                                                                                   (first entry)
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98US-0085295P.
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                                                            immunoglobulin
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Pred. No. 4.2e-11;
Mismatches 0;
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                                                            mite; antiallergic; mite; I
lobulin E; hypersensitivity;
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Best Local S
Matches 19
04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated mite allergenic protein of Dermatophagoides, designated Der HMW-map protein, and its related nucleic acid. The Der HMW-map protein is useful for eliciting an immune response against Der HMW-map protein. The protein or a reagent comprising a non-proteinaceous epitope is useful for identifying an animal (e.g., dog, cat) susceptible to or having an allergic response to a mite. A therapeutic composition is useful for desensitising a host animal to an allergic response to a mite. The DNA and protein can be used in the detection of anti-Der HMW-map antibodies in animal fluids, and inhibition of immunoglobulin (Ig)E or Der HMW-map protein activity associated with a disease. Antibodies that bind to Der HMW-map are useful for inhibiting being the force of the protein activity associated with a classes. Antibodies that bind to Der HMW-map are useful for inhibiting being the force of the protein activity associated with a classes.
                                                                                                                                                           30-JAN-2001; 2001WO-US000670
                                                                                                                                                                                                           09-AUG-2001.
                                                                                                                                                                                                                                                               WO200157278-A2
                                                                                                                                                                                                                                                                                                                                                                                          Probe; human; microarray; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAM18898;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               binding of proteins to IgE, to prevent immunocomplex formation, thus reducing hypersensitivity responses to mite allergens, and as vaccine against mite allergen hypersensitivity. Sequences AdU96314-AAU96342 represent Der HMW-map polypeptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New mite allergenic protein isolated from Dermatophagoides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-351888/38
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; protein; 20
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2000US-0180312P.
2000US-0207456P.
2000US-00608408.
2000US-00632366.
2000US-0234687P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                             encoded by probe for measuring cervical gene expression
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Pred. No. 4.2e-11;
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                                                                                                                                                                                                                                                                                                                                                                                            expression; cervical epithelial cell;
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Best Local
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26-MAY-2000;
30-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-SEP-2000;
04-OCT-2000;
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                             Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver.
                                                                                                                                                  Penn
                                                                                                                                                                                                                                            04-OCT-2000;
                                                                                                                                                                                                                                                                                                                    03-AUG-2000;
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                                                                                                                                                                                               MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LPHP-HSHTHLLCQ 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 23724; 487pp; English
                                                                                                                                                Hanzel
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2000US-00632366.
2000US-006323667P.
2000US-0236359P.
2000US-0024263.
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2000GB-00024263.
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Pred. No. 58;
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le encoded
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expression

Best Local Similarity

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RESULT 7
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Best Local
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26-MAY-2000;
30-JUN-2000;
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21-SEP-2000;
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                                                                                                                                       The present invention relates to single exon nucleic acid probes (SENP: see ANI3)315-AAI57546). The present sequence is a peptide encoded by on such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of the probes are useful for antenatal diagnosis of
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                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                         Human genome-derived single exon nucleic acid gene expression in human placenta.
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                                                                                                                     genetic disorders
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34.4%;
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Pred. No. 58;
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No. 58;
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                                                                                                                  The present invention relates to single exon nucleic acid probes for measuring human gene expression in a sample derived from human heart (see ABA21335-ABA41305). The present sequence is a protein encoded by one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                          04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
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2000US-00632366.
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2000US-0236359P.
2000US-0024263.
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Pred. No. 58;
5; Mismatches
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RESULT

WO200157275-A2

Homo

Alzheimer's disease;

Human; brain expressed exon; gene expression analysis; probe; microarray;

multiple sclerosis;

schizophrenia; epilepsy; cancer

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                                                                                                                                                          Human brain expressed single
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26-MAY-2000; 2000US-0207456P.
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03-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                          exon probe encoded protein SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probes useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                          NO: 30752
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RESULT 1:
ABG52865
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Best Local S
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03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                           03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                       04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                    probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is a protein encoded by one of the probes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Penn
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26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-AUG-2001.
                                                                                                 30-JAN-2001; 2001WO-US000664
                                                                                                                     09-AUG-2001
                                                                                                                                        WO200157273-A2
                                                                                                                                                          Homo sapiens
                                                                                                                                                                              hypercholesterolaemia;
                                                                                                                                                                                        Human; liver;
                                                                                                                                                                                                           Human liver peptide,
                                                                                                                                                                                                                               25-FEB-2003
                                                                                                                                                                                                                                                                     ABG52865 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       brains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
 (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MOLE-)
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 MOLECULAR DYNAMICS INC
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                                                                                                                                                                                                                                                                                                                     LPHP-HSHTHLLCQ 20
                                                                                                                                                                                                                                                                                                                                        IPHPTNIHKYLVCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hanzel DK,
                 2000US-0180312P.
2000US-0207456P.
2000US-0060840B.
2000US-00632366.
2000US-0234687P.
2000US-0236359P.
2000US-0236359P.
2000US-0236359P.
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2000US-0207456P.
2000US-0060840B.
2000US-00632366.
2000US-0234687P.
2000US-0236359P.
2000US-0236359P.
2000US-0236359P.
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K
                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                              (first
                                                                                                                                                                              cirrhosis; hyperlipoproteinaemia; hyperlipidaemia;
rolaemia; coronary heart disease.
                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                    34.4%;
                                                                                                                                                                                                           SEQ ID No 31513.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30752;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probes
                                                                                                                                                                                                                                                                                                                                                           Score 37.5; D. Pred. No. 58; 5; Mismatches
                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                      Å
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rank DR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for analyzing gene expression in human
                                                                                                                                                                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                                                                                                                                                                               4.
                                                                                                                                                                                                                                                                                                                                                            2;
                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                               20;
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RESULT 12
ABG40950
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                      04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                          Human; single exon probe; asthma; lung cancer; COPD; ILD; chronic obstructive pulmonary disease; interstitial lung disease; familial idiopathic pulmonary fibrosis; neurofibromatosis; tuberous sclerosis; Gaucher's disease; Niemann-Pick disease; hermansky-pudlak syndrome; sarcoidosis; pulmonary haemosiderosis; pulmonary histiocytosis; lymphangioleiomyomtosis; Karagener syndrome; pulmonary alveolar proteinosis; fibrocystic pulmonary dysplasia; primary ciliary dyskinesis; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stringency to a nucleic acid molecule expressed in the human adult liver. (I) may be used for predicting, measuring and displaying gene expression in samples derived from human adult liver. The genes identified may be involved in genetic liver diseases such as cirrhosis, hyperlipoproteinaemia, hyperlipidaemia and hypercholesterolaemia which is associated with coronary heart disease. ABG47348-ABG5930 represent human liver single exon encoded peptides of the invention. Note: The sequence information for this patent does not appear in the printed specification but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                  Human peptide
                                                                                                                                                                                                                                                                                                                                                                                                               19-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABG40950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              liver, comprising one of 13109 defined nucleotide sequences given in specification (or complements/ fragments). The probe hybridises at his
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            measuring human gene expression in a
                                                                                                                                    30-JAN-2001;
                                                                                                                                                                15-NOV-2001
                                                                                                                                                                                                                                                hyaline membrane
 (MOLE-)
                                                                                                                                                                                                                     sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           genome-derived single excepression in human adult
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 MOLECULAR DYNAMICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IPHPTNIHKYLVCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LPHP-HSHTHLLCQ 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA;
                        2000US-0207456P.
2000US-00638408.
2000US-00632366.
2000US-0234687P.
2000US-0236359P.
2000US-0236359P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                       2001WO-US000665
                                                                                                            2000US-0180312P
                                                                                                                                                                                                                                                                                                                                                                                    encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ID NO 31513; 658pp;
                                                                                                                                                                                                                                                disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34.4%;
                                                                                                                                                                                                                                                                                                                                                                                    by genome-derived single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to a single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         exon nucleic acid probes useful for analyzing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>ა</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 37.5;
Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rank
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            exon nucleic acid
n a sample derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2;
                                                                                                                                                                                                                                                                                                                                                                                      exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               de sequences given in the probe hybridises at high in the human adult liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            from human adult
                                                                                                                                                                                                                                                                                                                                                                                    probe
                                                                                                                                                                                                                                                                                                                                                                                      SEQ
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Compresents of the 1230, West reaching the movel set of probes

Compress Also inclided are a microarray comprising the novel set of probes

Compress of the cover set of probes which hybridise at high stringency to a nucleic

Collection of detectably labeled nucleic acids derived from human lung, measuring gene expression in a sample

ComRNA, and (b) measuring the label detectably bound to each probe of the

Comrany; identifying exons in a eukaryotic genome, comprising (a)

algorithmically predicting at least one exon from genomic sequences of

Compression of detectably hound to each probe of the

Comprising (a) identifying exons from genomic sequences of

Comprising (a) identifying exons from genomic sequences of

Comprising (a) identifying exons from genomic sequences of

Comprising (a) identifying exons from genomic sequence by the method

Comprising (a) identifying exons from genomic sequence by the method

Comprising (a) identifying exons from genomic sequence by the method

Comprising (a) identifying exons from genomic sequence by the method

Comprising (a) identifying exons from genomic sequence by the method

Comprising (a) identifying exons from genomic sequence by the method

Comprising (a) identifying exons from genomic sequence by the method

Comprising (b) measuring the expression of each of the exons in several

Comprising (b) measuring the expression of each of the exons in several

Comprising (c) identifying exons in a gene; a peptide comprising one

Comprising (c) identifying exons in a gene; a peptide comprising one

Comprising (c) identifying exons in a gene, particularly using human

Comprising (c) identifying exons in a gene, particularly using human

Comprising (c) identifying exons in a gene, particularly using human

Comprising (c) identifying exons in a gene, particularly using human

Comprising (c) identifying exons in a gene, particularly using human

Comprision (c) identifying exons in a gene, particularly using human

Comprision (c) identifying exons in a gene, particularly using hum
                                                                                                                                                                                                                                             RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
target stain; phenol oxidising enzyme; redox reaction; detergent industry; paper and pulp industry; textile industry; food industry; laccase enzyme; textile; carotenoid stain; fabric;
                                                                                                                             18-DEC-2003
                                                                                                                                                                                                       ADC27861 standard; peptide; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human lung comprising single exon nucleic acid probes having one 12614 nucleic acid sequences mentioned in the specification, or their complements or the 12387 open reading frames derived from the 12614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Spatially addressable set of single exon nucleic acid probes, used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Penn
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                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27; SEQ ID NO 30615;
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                                                                                                                                                                                                                                                                                                                                                                                         Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                IPHPTNIHKYLVCE 15
                                                                                                                                                                                                                                                                                                          LPHP-HSHTHLLCQ 20
                                                                                  stain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hanzel DK,
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                                                                                                                        (first entry)
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                                                                              targetting peptide Seq ID54
                                                                                                                                                                                                                                                                                                                                                                                                            34.4%;
42.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     634pp; English
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                                                                                                                                                                                                                                                                                                                                                                                       Score 37.5; D
Pred. No. 58;
5; Mismatches
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Matches
                                                      Query Match
                                                                                                                                                                                 This invention relates to novel peptides which bind to a target stain. The invention comprises phenol oxidising enzyme-binding peptide complexes wherein the binding peptide is attached to the C-terminus of the phenol oxidising enzyme or is inserted or substituted into the phenol oxidising enzyme. Phenol oxidising enzymes function by catalysis of redox reactions and are used for a wide range of applications (detergent industry, paper and pulp industry, textile industry and food industry). The invention is useful for enhancing the binding of a laccase enzyme to a target stain on a textile, where the peptide binds to the stain and not the textile. This may be useful for bleaching a carotenoid stain on a fabric or surface. The invention is useful in skin cosmetics and in food as fruit ripening agents. The present sequence is that of a stain targetting peptide of the
                                                                                                            Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 54; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       enzyme to a target stain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel binding peptide useful enzyme to a target stain, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-532545/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aehle W, Baldwin TL,
Wang H, Winetzky DS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-SEP-2002; 2002WO-US027993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           skin cosmetic; food; fruit ripening agent; oxidase protein; stain targetting peptide.
                                                                                                                                                                   invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-SEP-2001; 2001US-00954385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEMV ) GENENCOR INT
                            Local
Similarity 83.
                                                                                                            ΑĄ;
                         32.1%;
83.3%;
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                         Score 35; DB 7;
Pred. No. 2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for enhancing the target a stain on
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     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murray CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          binding of a laccase
a textile, and in skin
                                                      Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Van
  Indels
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-JUN-2005
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03-SEP-2002; 2002US-00235043 03-SEP-2002; 2002US-00235043

Synthetic US2005058996-A1.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC display peptides (ANYS6178-ANYS6609); to carotenoid-binding peptides (CC which comprise a repeatable motif selected from PPP, SSP, SSK, SPT, PLP, CC SLH, SPL, TTT, NTS, APS, TPV or a motif shown in ADYS663-ADYS6631; and CC stachybotrys laccase such as ADYS6177 or variants thereof) covalently a CC stachybotrys laccase such as ADYS6177 or variants thereof) covalently a CC also relates to polynucleotides encoding a carotenoid-binding peptide, CC and expression vectors and host cells comprising a polynucleotide encoding a carotenoid-binding peptide, CC and expression vectors and host cells comprising a polynucleotide encoding a phenol-oxidizing enzyme/carotenoid-binding peptide complex. CC condizing a phenol-oxidizing enzyme/carotenoid-binding peptide complex. CC condizing enzyme for enhancing the selectivity of the enzyme to a target carotenoid stain on a textile or other surface, where the enzyme can then CC condustry in the treatment, processing, finishing, polishing or production of fibers. The enzyme/peptide complexes are additionally useful in CC feach as in pharmaceutical applications e.g., to localize the presence of carotenoids in tissues. The present sequence represents a geolifically claimed carotenoid-binding cyclic peptide isolated from a CC charge display claimed carotenoid-binding cyclic peptide isolated from a conspectifically claimed carotenoid-binding cyclic peptide isolated from a consecution of the cyclic peptide isolated from a consecution of the cyclic peptide isolated from a consecution of the cyclic peptide isolated from a cyclic peptide isolated from a cyclic peptide isolated from a cyclic peptide cyclic peptide isolated from a cyclic peptide cyclic peptide isolated from a cyclic peptide cycli
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Best Local S
Matches
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Wang H,
                                                                                                                        cancer; neoplasm
                                                                                                                                           cytostatic; vaccine; gene therapy; pharmaceutical; immunotherapy; tumor;
                                                                                                                                                                                H3 or D3-acetylated tumor-associated peptide
                                                                                                                                                                                                                           20-OCT-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 54; 107pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptides that bind to carotenoid compounds, useful as detergents for targeting or removing stains on textiles or fabrics, in personal care or food industry applications, or in various diagnostic applications.
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(BALD/)
(VGAS/)
(JANS/)
  18-AUG-2005
                                        WO2005076009-A2
                                                                                                                                                                                                                                                                                                           AEC06055 standard;
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JANSSEN G G.
MURRAY C J.
WANG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 5; Conserv
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Winetzky DS;
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83.3%;
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Pred. No. 2e+0
1; Mismatches
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2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a method of identifying and quantifying tumoral sasociated peptides comprising chemically identically modifying peptides from samples in order to generate different physical characteristics in the peptides from the different samples. The peptides are useful for producing a medicament for the treatment of tumorous diseases and/or adenomatous diseases, such as renal, lung, colon, stomach, pancreatic, breast, prostate, ovarian and/or skin cancer. The peptide is used together with an adjuvant. A peptide bound to an antigen-presenting cell is used. The peptides are also useful for the labeling of leukocytes, in particular of T-lymphocytes, for evaluating the progress of therapy in a tumorous disease, and for producing an antibody. The nucleic acid and/or vector and/or cell are useful in producing a medicament for the treatment of tumorous and/or adenomatous diseases. This is the amino acid sequence of a tumor-associated peptide that binds human MHC class I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 5; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 10
                                                                         28-JAN-2004;
06-MAR-2004;
                                                                                                                                                                                     WO2005076009-A2
                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                              cytostatic; vaccine; gene therapy; pharmaceutical; immunotherapy; tumor;
                                                                                                                                                                                                                                                                                               Tumor-associated peptide SEQ ID NO 12.
                                                                                                                                                                                                                                                                                                                              20-OCT-2005
                                                                                                                                                                                                                                                                                                                                                                                        AEC06029 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying and quantifying tumor-associated peptides by chemically identically modifying peptides to generate different physical characteristics, useful for treating tumorous and/or adenomatous
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06-MAR-2004; 2004DE-10011503:
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                                                                                                                      28-JAN-2005; 2005WO-EP000873
                                                                                                                                                      18-AUG-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (IMMA-) IMMATICS BIOTECHNOLOGIES GMBH
                                           (IMMA-) IMMATICS BIOTECHNOLOGIES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                         2004DE-10005273
2004DE-10011503
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Pred. No. 70;
2; Mismatches
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Lemmel C,

Rammensee H

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RESULT 17
AAW30771
ID AAW30
XX AAW30
XX AAW30
XX Prost
PR O6-FE
XX Prost
PR WPI;
XX Prost
PT Cancr
XX Prost
PT Cancr
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes a method of identifying and quantifying tumorassociated peptides comprising chemically identically modifying peptides from samples in order to generate different physical characteristics in the peptides from the different samples. The peptides are useful for producing a medicament for the treatment of tumorous diseases and/or adenomatous diseases, such as renal, lung, colon, stomach, pancreatic, breast, prostate, ovarian and/or skin cancer. The peptide is used cogether with an adjuvant. A peptide bound to an antigen-presenting cell is used. The peptides are also useful for the labeling of leukocytes, in particular of T-lymphocytes, for evaluating the progress of therapy in a tumorous disease, and for producing an antibody. The nucleic acid and/or vector and/or adenomatous diseases. This is the amino acid sequence of a tumor-associated peptide that binds human MHC class I.
                                                                                                                             WPI; 1997-415352/38
                                                                                                                                                            Pettersson
                                                                                                                                                                                                                                        06-FEB-1996;
                                                                                                                                                                                                                                                                        06-FEB-1997;
                                                                                                                                                                                                                                                                                                       14-AUG-1997
                                                                                                                                                                                                                                                                                                                                     WO9729199-A2
                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                             Prostate specific antigen; PSA; immunogenic; human glandular kallikrein; hK2; antigen; antibody; detection; diagnosis; prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prostate Specific Antigen (PSA) derived peptide ABT3.
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                                                                                              Prostate Specific Antigen peptide(s)
                                                                                                                                                                                                         (ABBO ) ABBOTT LAB
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                                                                                                                                                            Bridon DP, IK;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry
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50.0%;
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Pred. No. 70;
2; Mismatches
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                                                                                              - useful
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70;
                                                                                                                                                                           Piironen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
                                                                                            for diagnosis of prostate
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AAW30769-84 are synthetic peptides antigen (PSA) sequence. These pepti

tides derived peptides are

from the pidentical

prostate specific l to a highly

Claim 2;

Page 10; 42pp; English.

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The present sequence represents a prostate specific antigen (PSA) (C) peptide. Monoclonal antibodies specific for PSA and hybridomas producing C them have been developed. The antibodies: (a) bind to free PSA; (b) are CC monoclonal antibody 365 binding to amino acids 82-87 of free PSA (LKNRFL) CC or fragment, and (c) are monoclonal antibodies 10, 11, 16 or 22.2 and C15.2, 156 or 225 binding to amino acids 139-144 (EELFIT) and 55-60 (SLFHPB) respectively of free and bound PSA, or fragments The antibodies CC comprising a solid support with attached labelled monoclonal antibody CC comprising a solid support with attached labelled monoclonal antibody CC comprising a solid support with attached labelled monoclonal antibody CC detect both free and bound PSA, a second solid support with attached (differently labelled) monoclonal antibody binding free and bound PSA (especially (b)) and a PSA standard can be used. To C (especially selected from (c)) can be used either with, or in place of, C (especially prostate and breast cancer. By obtaining total and free PSA values, their ratio can be used to separate prostatic cancer (PCa) from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                       Monoclonal antibodies specific for prostate specific antigen - useful, e.g. in screening for prostate or breast cancer and especially to distinguish between benign prostatic hyperplasia and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunogenic region of PSA, and also comprise one or more amino acids identical or non-identical to the amino acid sequence of human glandular kallikrein (hKZ). The peptides are used as antigens for the production of antibodies which are used to detect PSA in a test sample (claimed). This is useful for diagnosis of prostate cancer. The peptides enable the production of antisera necessary to determine the amount of total PSA, free PSA and PSA-ACT complex present in a sample and thus improve the ability of the clinician to distinguish, e.g., between BPH (benign
                                                                                                                                                                                                                                                                                                                                     Example; Page 59; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-193789/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Heavner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-SEP-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; prostate specific antigen; PSA; epitope; monoclonal antibody; detection; cancer; serine protease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CENZ ) CENTOCOR INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human prostate specific antigen peptide SEQ ID NO:72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW58054 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 35; DB
Pred. No. 1.1e
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e 35; DB 2; Lex.
No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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them have been developed. The antibodies: (a) bind to free PSA; (b) are monoclonal antibody 365 binding to amino acids 82-87 of free PSA; (b) are 15.2, 156 or 225 binding to amino acids 82-87 of free PSA (LKNRFL) cor fragment, and (c) are monoclonal antibodies 10, 11, 16 or 22.2 and 15.2, 156 or 225 binding to amino acids 139-144 (EELPLIP) and 55-60 (SIFHPE) respectively of free and bound PSA, or fragments. The antibodies are useful to detect PSA. For detecting free PSA only, an immunoassay comprising a solid support with attached labelled monoclonal antibody specific for free PSA (especially (b)) and a PSA standard can be used. To detect both free and bound PSA, a second solid support with attached (differently labelled) monoclonal antibody binding free and bound PSA (especially selected from (c)) can be used either with, or in place of, the first solid support. The antibodies are useful in cancer screening, especially prostate and breast cancer. By obtaining total and free PSA values, their ratio can be used to separate prostatic cancer (PCa) from the support of the static cancer. By obtaining total and free PSA is also useful after radical prostatectomy, to predict disease persistence. The antibodies allow evaluation of PSA free/total ratio, enabling separation of PSA free/total ratio, enabling separation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monoclonal antibodies specific for prostate specific antigen - useful e.g. in screening for prostate or breast cancer and especially to distinguish between benign prostatic hyperplasia and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15
antibodies of BPH and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   peptide. Monocional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Page 60; 84pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-193789/17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Heavner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9810292-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detection; cancer; serine protease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; prostate specific antigen; PSA; epitope; monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW58055
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CENZ ) CENTOCOR INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present sequence represents a prostate specific antigen (PSA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           prostatic hyperplasia (BHP) patients. Measurement of PSA is after radical prostatectomy, to predict disease persistence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HPTNIHKYLVC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HPOKVTKFMLC 15
   PCa
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patients with PSA values 4-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96US-0025404P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US014909
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibodies specific for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        32.1%;
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Pred.
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No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PSA and
ng/ml not
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Query Match Best Local Matches

4: Similarity

32.1%; milarity 36.4%; Conservative

4.

Score 35; DB 7; Pred. No. 1.1e+02; 4; Mismatches 3

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DB 7;

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RESULT 20
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                                                 The invention relates to a composition comprising at least one peptide CC having an isolated, prepared epicope selected from any of the sequences CC from 30 lists given in the specification. Also disclosed is a method for CC inducing a cytotoxic T cell response against a pre-selected antigen in a CC patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The CC composition comprises an epitope that is joined by an amino acid linker. CC The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an A2 crestricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL chinds to a complex of the HLA molecule and the epitope. Specifically CC claimed are peptides having allele-specific binding motifs for HLA. The CC compositions and methods are useful for preventing, treating or CC diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADM29251-CC ADM37745 represent epitopes of the invention as given in Tables 2-31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             possible by total PSA testing, avoiding biopsies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-MAR-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            binding motifs for HLA, us
viral diseases and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sette A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-OCT-2000; 2000US-0242350P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              viral disease; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HLA binding
   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 52-379; 382pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-441519/41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sidney J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,
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Pred. No. 1.1e+02;
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                                                                                                                              RESULT 21
ADW35209
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                                                             Best Loc
Matches
                                                                                             Query Match
                                                                                                                                                        claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnosing viral diseases and cancer. The peptide epitopes are useful as diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
                                                                                                                                                                                                                                                         composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an Alrestricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing viral diseases and cancer.
                                                                                                                                                                                                                                                                                                                                         The invention relates to a composition comprising at least one peptide having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The
                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 52-379; 382pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T cell; MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA; viral disease; cancer.
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20-APR-2001;
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                                                                                Local
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                                                             Similarity
4; Conserv
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HPQKVTKFMLC 15
                             HPTNIHKYLVC 14
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                                                               Conservative
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2001US-0285624P.
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                                                                              36.4%;
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                                                             Score 35; DB 7; Le
Pred. No. 1.1e+02;
4; Mismatches 3;
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                                                                                             Length 15
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                                                           Gaps
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RESULT

22

10-MAR-2005

(first entry)

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RESULT 23
ADW33976
ID ADW33
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AC ADW33
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AC ADW33
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DT 10-M2
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             patient expressing a specific MHC class I allele by contacting cyctoxic T cells from the patient with the composition cited above. The composition comprises an epitope that is joined by an amino acid linker. The epitope is admixed or joined to a CTL or HTL epitope. The epitope is bound to an HLA molecule on the antigen-presenting cell, where when an A2 restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically claimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW29251-ADW37745 represent epitopes of the invention as given in Tables 2-31.
                                                                                        ADW33976;
                                                                                                                                           ADW33976 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing viral diseases and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a composition comprising at least one peptide having an isolated, prepared epitope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method foi inducing a cytotoxic T cell response against a pre-selected antigen in a inducing a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in a cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pre-selected antigen in the cytotoxic T cell response against a pr
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20-APR-2001; 2001US-0285624P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T cell; MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADW33268 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPIM-) EPIMMUNE INC
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                                                                                                                                                                                                                                                                                                                                                            4 HPTNIHKYLVC 14
                                                                                                                                                                                                                                                                                                                                                                                                                           4; Conservative
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HPQKVTKFMLC
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                                                                                                                                                                                                                                                                                                                                                                                                                        4; Mismatches
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Pred. No. 1.1e+02;
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RESULT 24
ADN65435
ID ADN65
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AC ADN65
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AC ADN65
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DT 01-JU
DT 01-JU
DX
DE HLA h
XX
GYTOS
KW CYTOS
KW GYTOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a composition comprising at least one peptide CC having an isolated, prepared epicope selected from any of the sequences from 30 lists given in the specification. Also disclosed is a method for inducing a cytotoxic T cell response against a pre-selected antigen in a patient expressing a specific MHC class I allele by contacting cytotoxic T cells from the patient with the composition cited above. The composition comprises an epicope that is joined by an amino acid linker. The epicope is admixed or joined to a CTL or HTL epicope. The epitope is composition comprises an epicope from the price of the CTL or HTL epicope. The epitope is domixed or joined to a CTL or HTL epitope. The epitope is compositione and had molecule on the antigen-presenting cell, where when an A2 restricted cytotoxic lymphocyte (CTL) is present, a receptor of the CTL binds to a complex of the HLA molecule and the epitope. Specifically CC laimed are peptides having allele-specific binding motifs for HLA. The compositions and methods are useful for preventing, treating or diagnosing viral diseases and cancer. The peptide epitopes are useful as C diagnostic agents for evaluating immune responses, for making antibodies and for evaluating efficacy of a vaccine. Sequences given in ADW39251- ADW37745 represent epitopes of the invention as given in Tables 2-31.
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                              Matches
cytostatic; hepatotropic; virucide; antiinflammatory; anti-HIV; gene therapy; vaccine; HIA binding peptide; HITL epitope; liposome; prostate specific antigen; prostate specific membrane antigen; hepatitis B virus antigen; hepatitis C virus antigen; malignant melanoma antigen; MAGE; Epstein Barr virus; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Virucide; cytostatic; gene therapy; vaccine; epitope; cytotoxic T cell; MHC class I; CTL; HTL; A2-restricted cytotoxic lymphocyte; HLA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New composition comprising at least one peptide having allele-specific binding motifs for HLA, useful for preventing, treating or diagnosing viral diseases and cancer.
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                                                                                                                    HLA binding peptide #2035
                                                                                                                                                            01-JUL-2004
                                                                                                                                                                                                  ADN65435;
                                                                                                                                                                                                                                      ADN65435 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15
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20-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HLA binding epitope #4726
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                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                    HPQKVTKFMLC 15
                                                                                                                                                                                                                                                                                                                                                                        HPTNIHKYLVC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sidney J,
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                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                          (first entry
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2001US-0285624P
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                                                                                                                                                                                                                                                                                                                                                                                                            Score 35; DB Pred. No. 1.1e
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. 1.1e+02;
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The invention relates to a novel composition comprising one or more CC composition further comprises an HTL epitope. It also comprises a spacer CC composition further comprises an HTL epitope. It also comprises a spacer CC molecule, a carrier, an MHC targeting sequence or a lipid. The peptides selected from prostate specific antigen (PSA), prostate specific membrane CC antigen (PSA), hepatitis B virus (HBV) antigen, hepatitis C virus (HCV) CC antigen, malignant melanoma antigen (MAGE), Epstein Barr virus, human CC immunodeficiency type-1 (HIV-I), human immunodeficiency type-2 (HIV-2), CC Papilloma virus, Lassa virus, Mycobacterium tuberculosis (WTV), p53, CCC murine p53 (mp53), CEA, HER2/neu, and tyrosine kinase related protein CC infections or cancer, such as prostate cancer, hepatitis B, hepatitis C, CC AIDS, renal carcinoma, cervical carcinoma, lymphoma, CMV or chondyloma CC acuminatum. The composition is also be used for diagnosing such diseases. Tris sequence represents a peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New composition of peptides and nucleic acids capable of binding Major Histocompatibility Complex molecules, useful for diagnosing, preventing or treating viral infections or cancer, such as prostate cancer, hepatitis B or AIDS.
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08-OCT-2002; 2002US-0417269P.
Sequence 15 AA;
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Best Local
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                                     Similarity
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              HPTNIHKYLVC 14
HPQKVTKFMLC
                              Conservative
                                     32.1%;
15
                                     Score 35; DB 8;
Pred. No. 1.1e+02;
                             Mismatches
                                             Length 15;
                              <u>.</u>
                              Gaps
                              0
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RESULT 25
AAW30783
ID AAW30
XX AAW30
XX AAW30
XX Prost
DT 27-FE
DT 27-FE
DX Prost
XX WProst
XX WProst
XX WProst
XX WW977
                                                                                                                                                                                  AAW30783
                                                                                                            Prostate
                                                                                                                                    27-FEB-1998
                                                                                                          Specific Antigen
                                                                                                                                                                                   standard;
                                                                                                                                  (first entry)
                                                                                                                                                                                  peptide; 19
                                                                                                            (PSA)
                                                                                                            derived peptide ABT15
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Prostate specific antigen; antibody; detection; PSA; immunogenic; human glandular kallikrein; diagnosis; prostate cancer

hK2; antigen;

Synthetic Homo sapiens

WO9729199-A2

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Search completed: January 26, 2006, 07:58:09 Job time : 68.5345 secs
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Matches 4; Conserv
                                                                                                                                                                                                                AAW30769-84 are synthetic peptides derived from the prostate specific antigen (PSA) sequence. These peptides are identical to a highly immunogenic region of PSA, and also comprise one or more amino acids identical or non-identical to the amino acid sequence of human glandular kallikrein (hK2). The peptides are used as antigens for the production of antibodies which are used to detect PSA in a test sample (claimed). This is useful for diagnosis of prostate cancer. The peptides enable the production of antisera necessary to determine the amount of total PSA, free PSA and PSA-ACT complex present in a sample and thus improve the ability of the clinician to distinguish, e.g., between BPH (benign prostatic hyperplasia) and prostatic cancer in a patient
                                                                                                                                                                                            Sequence 19 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 10; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prostate Specific Antigen peptide(s) - useful for diagnosis of prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-415352/38.
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Pred. No. 1.5e+02;
4; Mismatches 3; Indels
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US-08-595-945-15
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US-09-926-276-20
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 11, Appli 3, Appli 20, Appli 21, App
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US-08-141-935-75
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ALIGNMENTS

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TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PROTEINS
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: AL-2-C3
CURRENT APPLICATION NUMBER: US/09/292,225
CURRENT FILING DATE: 1999-04-15
EARLIER FILING DATE: 1999-09-02
EARLIER FILING DATE: 1998-05-13
EARLIER FILING DATE: 1998-05-13
EARLIER APPLICATION NUMBER: 60/098,565
EARLIER APPLICATION NUMBER: 60/098,565
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EARLIER FILING DATE: 1998-04-17
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SOFTWARE: PATENTON SEQ ID NOS: 49
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SOFTWARE: PATENTON SEQ ID NOS: 49
SOFTWARE: Dermatophagoides farinae
US-09-292-225-11
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                                                                   SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
LENGTH: 20
TYPE: PRT
ORGANISM: Dermatophagoides farinae
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Patent No. 6455686
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APPLICANT: Hunter, Shirley Wu
APPLICANT: Weber, Eric R.
APPLICANT: Weber, Eric R.
TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PROTEINS
TITLE OF INVENTION: AND USES THEREOF
FILE REPERENCE: AL-2-C3
CURRENT APPLICATION NUMBER: US/09/292,225
CURRENT FILING DATE: 1999-04-15
EARLIER APPLICATION NUMBER: 60/098,909
EARLIER FILING DATE: 1998-09-02
EARLIER FILING DATE: 1998-09-02
                                                                                                                                                                                                  EARLIER APPLICATION NUMBER: 60/085,295
EARLIER FILING DATE: 1998-05-13
EARLIER APPLICATION NUMBER: 60/098,565
EARLIER FILING DATE: 1998-04-17
EARLIER APPLICATION NUMBER: 09/062,013
EARLIER FILING DATE: 1998-04-17
NUMBER OF SEQ ID NOS: 49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ch 100.0%; Score 109; DB 2; 1 Similarity 100.0%; Pred. No. 2.6e-11; 19; Conservative 0; Mismatches 0;
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  Score 109;
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Length 20;
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US-08-595-945-15

Sequence 15, Application US/08595945
Patent No. 6143509
GENERAL INFORMATION:

APPLICANT: APPLICANT:

DOWELL, BARRY L.
BRIDON, DOMINIQUE P.
QIU, XIAOXING
LILJA, HANS

APPLICANT:

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RESULT 4
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                                                                                                      Matches
                                                                                                               Query Match
Best Local Similarity
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Versi
RAPPLICATION NUMBER: US/08/595,945
FILING DATE: 06-FEB-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: BECKER, CHERYL L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 5875.US.01
                                                                                                                                                                                                                                                                      TELEFAX: 847-938-2623
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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APPLICANT:
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ADDRESSEE: ABBOTT LABORATORIES
STREET: 100 ABBOTT PARK ROAD
                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: PROSTATE SPECIFIC ANTIGEN PEPTIDES TITLE OF INVENTION: AND USES THEREOF
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                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                       LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                                                      4 HPTNIHKYLVC 14
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                                          HPOKVTKFMLC 15
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                                                                                                 32.1%;
ilarity 36.4%;
Conservative
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BRIDON, DOMINIQUE
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Pred. No. 33;
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APPLICANT: Kenten, John H
APPLICANT: Tramontano, Alfonso
APPLICANT: Tramontano, Alfonso
APPLICANT: Pilon, Aprile L
APPLICANT: Cohnas, Gerald L
APPLICANT: Roberts, Steven F
ITITE OF INVENTION: HEAT-SHOCK FUSION-BASED VACCINE SYSTEM
FILE REFERENCE: U.S. Patent Application No. 6319503 09\026,276
CURRENT APPLICATION NUMBER: US/09/026,276
CURRENT FILING DATE: 1998-02-19
NUMBER OF SEQ ID NOS: 35
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 20
LENGTH: 20
TYPE: PRT
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                                                                                                                                        ; ORGANISM: Homo sapiens
US-09-026-276-20
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SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
TYPE: amino acid
TYPE: amino acid
TYPE: amino acid
TYPE: 11near
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Best Local Similarity
Matches 4; Conserv
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OPERATING SYSTEM: PC-DOS/MS-DOS

SOPTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/595,945

FILING DATE: 06-FEB-1996

CLASSIFICATION: 435

ATTORNEY/AGENT INFORWATION:

NAME: BECKER, CHERYL L.

REGISTRATION NUMBER: 35,441

REFERENCE/DOCKET NUMBER: 35,411

REFERENCE/DOCKET NUMBER: 35,411

REFERENCE/DOCKET NUMBER: 35,411

REFERENCE/DOCKET NUMBER: 37,411

REFERENCE/DOCKET NUMBER: 37,411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Patent No. 6319503
                                                                    Query Match
                                   Best Local Similarity
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: ABBOTT LABORATORIES
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
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APPLICANT: VIHIMEN, MAUNO A.
APPLICANT: PETTERSSON, KIM S.I.
TITLE OF INVENTION: PROSTATE SPECIFIC ANTIGEN PEPTIDES
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 16
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   Conservative
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                                   32.1%; Score 35; DB 36.4%; Pred. No. 45;
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   Mismatches
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APPLICANT: Pilon, Aprile L
APPLICANT: Lohnas, Gerald L
APPLICANT: Lohnas, Gerald L
APPLICANT: Roberts, Steven F
APPLICANT: Roberts, Steven F
ITILE OF INVENTION: HEAT-SHOCK FUSION-BASED VAC
FILE REFERENCE: U.S. Patent Application No. 66/
CURRENT APPLICATION NUMBER: US/09/964,201A
CURRENT FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 20
LENGTH: 20
TYPE: PRT
CURRENT HOMO Sapiens
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US-07-829-462-6
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TELEFAX: 619-535-8949
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Ruoslahti, l
APPLICANT: Morla, Alex
                              ATTORNEY/AGENT INFORMATION:
NAME: CAMPBELL, CATHRYN A.
REGISTRATION NUMBER: 31,547
REFERENCE/DOCKET NUMBER: P-L.
TELECOMMUNICATION INFORMATION:
TELECHINE: 619-35-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Kenten, John APPLICANT: Tramontano,
                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: TITLE OF INVENTION:
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Local Similarity 36.4%;
                                                                                                                                                   APPLICATION NUMBER: US
FILING DATE: 19920131
CLASSIFICATION: 530
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T: 4370 La Jolla Village Drive, Suite 700
San Diego
California
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Pilon, Aprile
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                  619-535-8949
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Alfonso
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Pred. No. 45;
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US-08-159-339A-551
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                                                                     Query Match
Best Local (
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APPLICANT: Kubo,
                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                   NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REPERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-WAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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ADDRESSEE: Townsend and Townsend and Crew LLP
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                                                                                                                                             STRANDEDNESS: SIN
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                                                                                                                                                                            TYPE: amino acids
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FILING DATE: 29-NOV
CLASSIFICATION: 424
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Local Similarity 45.5%;
es 5; Conservative
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5; Conserv
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IPHPAGLKK 10
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                                                        Score 32; DB Pred. No. 66; 1; Mismatches
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US-09-009-953-109
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US-09-962-756-51
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Patent No. 6413517
GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
                                                                                                                                   GENERAL INFORMATION:
APPLICANT: PILLUTLA, RENUKA
APPLICANT: BRISSETTE, RENE
APPLICANT: BLUME, ARTHUR J
                                                                                                                                                                                                                Sequence 51, Application US/09962756 Patent No. 6875741
                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity
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                                                            APPLICANT:
                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: <Unknown:
INFORMATION FOR SEQ ID NO: 109:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 60/036,713
FILING DATE: 23-JAN-1997
APPLICATION NUMBER: US 60/037,432
FILING DATE: 07-FEB-197
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-011520US
CANT: BRANDT, JAKOB
CANT: GOLDSTEIN, NEIL I.
CANT: SPETZLER, JANE
CANT: OSTERGAARD, SOREN
CANT: HANSEN, PER HERTZ
CANT: HANSEN, PER HERTZ
CANT: HANSEN, PER HERTZ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 109:
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SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
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ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Identification of Broadly
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FILING DATE: 21-Jan-1998
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                     IPHPAGLKK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
                                                                                                                 BRISSETTE, RENEE
BLUME, ARTHUR J.
SCHAFFER, LAUGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 32; DB 2;
Pred. No. 1e+02;
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SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 51
LENGTH: 21
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CURRENT APPLICATION NUMBER: US/09/962,756
CURRENT FILING DATE: 2001-09-24
PRIOR APPLICATION NUMBER: 09/538,038
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
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                                                                                                       TELEFAX: 617-345-9111 NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                  APPLICATION NUMBER: US08, FILING DATE: 01-SEP-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: Wordperfect 6.1 CURRENT APPLICATION DATA:
 MOLECULE TY
                                                                                                                                         REFERENCE/DOCKET NUMBER: 32
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-9100
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Masucci,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: SE9501324-9 FILING DATE: 10-APR-1995
                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08 FILING DATE: 15-SEP-1995 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: One I
                                 STRANDEDNESS:
                                                                                                                                                                                          NAME: Williams, Ph.D., Kathleen A REGISTRATION NUMBER: 34,380
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                                                                     ENGTH:
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                                                 amino acid
TYPE:
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                unknown
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                                 single
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37.5%; Pred. No. 1.
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GLYCINE-CONTAINING SEQUENCES
CONFERRING INVISIBILITY TO T
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US-09-791-524A-3
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Patent No. 565672
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Best Local Similarity
Matches 5; Conserv
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/IB99/01524
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/098,028
PRIOR FILING DATE: 1998-08-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/791,524A CURRENT FILING DATE: 2001-02-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.2
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TITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
                              CLASSIFICATION: 530 PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH: 16
                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Doyle, Michael
APPLICANT: Goodson, Robert
TITLE OF INVENTION: Peptide Inhibitors of Urokinase Receptor
TITLE OF INVENTION: Activity
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                       STREET: 4500
CITY: Emeryville
                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                  APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                        MEDIUM TYPE:
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                 FILING DATE
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5; Conserv
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                                                                                                                                                                                                                                                                                             4560 Horton Street
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                                                                                                                                                                                       Floppy disk
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45.5%;
                                  US/08/061,514
                                                                                                  US/08/370,567
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Pred. No. 1.6e+02;
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Pred. No. 4.6e+05
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Patent No. 5679782
GENERAL INFORMATION:
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Best Local Similarity
Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO:
                                                                                                                                  TELEPHONE: 510-601-2706
TELEPAX: 510-655-3542
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
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LENGTH: 17 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Rosenberg, Steven
APPLICANT: Doyle, Michael
APPLICANT: Goodson, Robert
TITLE OF INVENTION: Peptide Inhibitors of Urokinase Receptor
TITLE OF INVENTION: Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: 510-601-2706
TELEFAX: 510-655-3542
                                                                   MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 510-601-2706
                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION: NAME: Green, Grant D.
                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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ADDRESSEE: Chiron Corporation
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                                                                                                                                                                                                                                        NAME: Green, Grant D.
REGISTRATION NUMBER: 31,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                   FILING DATE:
APPLICATION NUMBER: US/08/061,514
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                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                               FILING DATE
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45.5%;
28.4%;
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Pred. No. 1.7e+02
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 Score 31; DB 1;
Pred. No. 1.7e+02;
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US-08-538-911-20
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US-09-402-641-10
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                          Sequence 10, Application US/09402641
PATENT NO. 6528619
GENERAL INFORMATION:
APPLICANT: BUERGLE, Markus
APPLICANT: GRAEFF, Heinrich
APPLICANT: KESSLER, Horrst
APPLICANT: KESSLER, UKTST
APPLICANT: KESSLER, Viktor Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Ve CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/538 and FILING DATE: CLASSIFTO:
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Matches 5; Conserv
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INFORMATION FOR SEQ ID NO:
APPLICANT:
              APPLICANT:
                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 1i:
MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Green, Grant D.
REGISTRATION NUMBER: 31,259
REFERENCE/DOCKET NUMBER: 04
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/069,352
FILING DATE:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Chiron Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Doyle, Michael
APPLICANT: Winter, Jill
TITLE OF INVENTION: Method For Selection Of Biologically
TITLE OF INVENTION: Active Peptide Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 17 amino acids
TYPE: amino acid
STRANDEDNESS: single
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o. 5750344
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KOPPITZ, Marcus
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NO
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Pred. No. 1.7e+02;
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US-09-411-067C-2
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                                              Sequence 196, Application US/09428082B
PATENT NO. 6660843
GENERAL INFORMATION:
APPLICANT: FEIGE, ULRICH
APPLICANT: LIU, CHUAN-FA
APPLICANT: CHESTHAM, JANET C.
APPLICANT: BOONE, THOWAS CHARLES
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: NUVAS, LLC
APPLICANT: HOUSTON, L.L.
TITLE OF INVENTION: USE OF A CONTEXT-DEPENDENT FUNCTIONAL ENTITY TO ENHANCE THE EFFI
TITLE OF INVENTION: AN AGENT
FILE REFERENCE: NUVAS1140
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Best Local Similarity
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                                                                                                                                                                                            ·09-428-082B-196
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CURRENT FILLING DATE: 2000-01-10
PRIOR APPLICATION NUMBER: EP97106024.9
PRIOR FILING DATE: 1997-04-11
PRIOR APPLICATION NUMBER: PCT/EP98/02178
PRIOR FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 10
                               FILE REFERENCE: A-527
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TITLE OF INVENTION: INHIBITORS FOR UROKINASE RECEPTOR
FILB REFERENCE: Case 20367US
                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 17
TYPE: PRT
ORGANISM: Homo sapiens
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APPLICATION NUMBER: US/09/428,082B FILING DATE: 1999-10-22
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SCHMITT, Manfred
WEIDLE, Ulrich
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Pred. No. 1.7e+02;
2; Mismatches
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Pred. No. 1.7e+02;
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; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: peptide ligand
US-09-155-260C-16
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US-09-155-260C-16
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LENGTH: 17
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Matches 5; Conservative
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                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Riemer, Christoph
APPLICANT: Burgle, Markus
TITLE OF INVENTION: Inhibitors for Urokinase
FILE REFERENCE: 100564-09040
CURRENT APPLICATION NUMBER: US/09/402,464
CURRENT FILING DATE: 2000-01-07
                                                                                                                                                         APPLICANT:
                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 08/623,361
PRIOR FILING DATE: 1996-03-28
NUMBER OF SEQ ID NOS: 76
SOFTWARE: Patentin Ver. 2.1
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PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1133
                                                                                                                                                    APPLICANT: Kessler, Horst
APPLICANT: Graeff, Heinrich
APPLICANT: Schmitt, Manfred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: CHAPMAN, HAROLD
TITLE OF INVENTION: PEPTIDE LIGANDS OF THE UROKINASE RECEPTOR
FILE REFERENCE: 014024-0284102
CURRENT APPLICATION NUMBER: US/09/155,260C
CURRENT FILING DATE: 1998-09-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/US97/05199
PRIOR FILING DATE: 1997-03-28
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                                                                                                               Magdolen, Viktor Wilhelm, Olaf G.
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Pred. No. 1.7e+02;
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; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 17
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-402-464-1
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US-09-791-524A-4
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PCT-US94-05591-20
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Matches
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                   Sequence 20, Application PC/TUS9405591
GENERAL INFORMATION:
APPLICANT: Doyle, Michael V.
TITLE OF INVENTION: Method For Selection Of Biologically
TITLE OF INVENTION: Active Peptide Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 45.5%;
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Aventis Pharmaceuticals Products Inc.
TITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
FILE REFERENCE: p26,992-B USA
CURRENT APPLICATION NUMBER: US/09/791,524A
CURRENT FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: PCT/IB99/01524
PRIOR FILING DATE: 1999-08-27
PRIOR FILING DATE: 1999-08-27
PRIOR FILING DATE: 1998-08-27
NUMBER OF SEQ ID NOS: 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: EP 97 106 024.9 PRIOR FILING DATE: 1997-04-11 NUMBER OF SEQ ID NOS: 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 17
TYPE: PRT
ORGANISM: Adenovirus
               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05591
                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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CITY: Emeryville
STATE: CA
                                                                                                                                                 ZIP: 94608
                                                                                                                                                                 COUNTRY:
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19-MAY-1994
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45.5%;
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Pred. No. 1.7e+02;
2; Mismatches 4
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INFORMATION FOR SEQ ID NO:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: PCT/US94/05684
FILING DATE: 19 MAY 1994
CLASSIFICATION:
                                                                                                                                                                                                                                                  NAME: Green, Grant D.
REGISTRATION NUMBER: 31,259
REFERENCE/DOCKET NUMBER: 099
TELECOMMUNICATION INFORMATION:
TELEPHONE: 510-601-2706
                                                                                                                         MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Chiron Corporation
TITLE OF INVENTION: Peptide Inhibitors of Urokinase Receptor
TITLE OF INVENTION: Activity
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hes 5; Conservative
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REGISTRATION NUMBER: 31
REFERENCE/DOCKET NUMBER:
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CITY: Emeryville
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TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                       TYPE: amino acid
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2 IPHPTNIHKYL 12 :|| | :||
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                                             Similarity 5; Conserv
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                                                             Score 31; DB 4;
Pred. No. 1.7e+02;
                                                Mismatches
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Aventis Pharmaceuticals Products Inc.
ITITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
FILE REFERENCE: P26,992-B USA
CURRENT APPLICATION NUMBER: US/09/791,524A
CURRENT FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: PC7/1899/01524
PRIOR FILING DATE: 1999-08-27
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/098,028
PRIOR FILING DATE: 1999-08-27
NUMBER OF SEQ ID NOS: 165
SOPTWARE: PATENTING DATE: 1998-08-27
NUMBER OF SEQ ID NOS: 165
SOPTWARE: PATENTING DATE: 1998-08-27
SEQ ID NO 9
LENGTH: 20
TYPE: PRT
ORGANISM: Adenovirus
US-09-791-524A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 25
US-09-791-524A-10
| Sequence 10, Application US/09791524A
| Patent No. 6911199
| GENERAL INFORMATION:
| APPLICANT: Aventis Pharmaceuticals Products Inc.
| TITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
| FILE REFERENCE: P26,992-B USA
| CURRENT APPLICATION NUMBER: US/09/791,524A
| CURRENT FILING DATE: 2001-02-22
| PRIOR APPLICATION NUMBER: US/09/791,524A
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Search completed: January 26, 2006, 08:07:03 Job time: 19.0201 secs
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US-09-791-524A-9
; Sequence 9, Application US/09791524A
; Patent No. 6911199
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; ORGANISM: Adenovirus
US-09-791-524A-10
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SEQ ID NO 10
LENGTH: 21
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Best Local Similarity 45.5%;
Matches 5; Conservative
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Perfect score:
Sequence:
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Listing first 100:
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Match
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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   DIPHPTNIHKYLVCESVNG
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Copyright (c) 1993 - 2006 Compugen
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US-10-218-743-11
US-10-218-743-3
US-09-864-761-38552
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US-10-912-512-54
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US-10-912-512-54
US-10-912-512-54
US-10-681-389-20
US-10-681-389-20
US-10-681-389-20
US-10-681-708-95
US-10-200-708-47
US-10-200-708-119
US-10-200-708-119
US-10-200-708-671
US-10-623-710-19
US-10-623-710-19
US-10-623-710-19
US-10-423-649A-9
US-11-008-653-128
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US-10-253-471-51
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               sequence
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11, Appl
38522, Appl
54, Appl
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65, Appl
620, Appl
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29, Appli
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ALIGNMENTS

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RESULT 2
US-10-218-743-3
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                                                                                                        PRIOR APPLICATION NUMBER: 60/098,909
PRIOR FILLING DATE: 1998-09-02
PRIOR PELICATION NUMBER: 60/085,295
PRIOR FILLING DATE: 1998-05-13
PRIOR APPLICATION NUMBER: 60/098,565
PRIOR FILLING DATE: 1998-04-17
PRIOR APPLICATION NUMBER: 09/062,013
PRIOR APPLICATION NUMBER: 09/062,013
PRIOR FILLING DATE: 1998-04-17
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PRIOR FILING DATE: 1998-05-13
PRIOR PELICATION NUMBER: 60/098,565
PRIOR FILING DATE: 1998-04-17
PRIOR APPLICATION NUMBER: 09/062,013
PRIOR TILING DATE: 1998-04-17
NUMBER OF SEQ ID NOS: 49
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LENGTH: 19
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/10218743
Publication No. US20030096779A1
GENERAL INFORMATION:
APPLICANT: McCall, Catherine A.
APPLICANT: Hunter, Shirley Wu
APPLICANT: Weber, Eric R.
                                           SOFTWARE:
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 11, Appropriate Publication No.
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                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: AL-2-C3
CURRENT APPLICATION NUMBER: US/10/218,743
CURRENT FILING DATE: 2002-08-13
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                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: US/09/292,225
PRIOR FILING DATE: 1999-04-15
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PRIOR APPLICATION NUMBER: US/09/292,225
PRIOR FILING DATE: 1999-04-15
PRIOR BENTICE: 1999-04-15
                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PROTEINS TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/218,743
CURRENT FILING DATE: 2002-08-13
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LENGTH: 20
TYPE: PRT
                                                                                            NUMBER OF SEQ ID NOS: 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: AL-2-C3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PROTEINS TITLE OF INVENTION: AND USES THEREOF
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FILING DATE: 1998-09-02
                                                                      PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           McCarr, Sur-
Hunter, Sur-
Ther, Eric R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DIPHPTNIHKYLVCESVNG 19
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US-09-864-761-38552
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
                                                                                                                                                                                                                  SEQ ID NO 38552
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Best Local
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                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers.
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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APPLICANT: Rank, David R.
APPLICANT: Hanzel, Bavid K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
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PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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FEATURE:
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.6
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL
                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 09/774,203
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PRIOR APPLICATION NUMBER: US 60/234,687
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                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                    LENGTH:
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les 19; Conser
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Pred. No. 1.2e-09;
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/954,385
CURRENT FILING DATE: 2001-09-12
NUMBER OF SEQ ID NOS: 433
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 54
LENGTH: 7
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Best Local Similarity
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APPLICANT: Baldwin, Toby L.
APPLICANT: Van Gastel, Franciscus J.C.
APPLICANT: Janssen, Giselle G.
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PRIOR APPLICATION NUMBER: US/09/954,385
PRIOR FILING DATE: 2001-09-12
NUMBER OF SEQ ID NOS: 433
                                                                 APPLICANT: Winetzky, Deborah S.
TITLE OF INVENTION: Binding Phenol Oxidizing Enzyme-peptide
TITLE OF INVENTION: Complexes
FILE REFERENCE: GC690
CURRENT APPLICATION NUMBER: US/10/912,512
CURRENT FILING DATE: 2004-08-05
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TITLE OF INVENTION: Binding Phenol Oxidizing
TITLE OF INVENTION: Complexes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: binding peptide
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R INFORMATION:
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Murray, Christopher J.
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Murray, Christopher J.
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EXPRESSED IN HELA, SIGNAL = 1.8
EXPRESSED IN HEART, SIGNAL = 1.9
EXPRESSED IN BRAIN, SIGNAL = 1.9
EXPRESSED IN ADULT LIVER, SIGNAL = 1.8
SWISSPROT HIT: P53820, EVALUE 8.80e+00
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42.9%;
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Pred. No. 1.7e+06;
1; Mismatches
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TITLE OF INVENTION: HEAT-SHOCK FUSION-BASED VACCINE SYSTEM FILE REFERENCE: U.S. Patent Application No. US20030091575A1 09\(CURRENT APPLICATION NUMBER: US/09/964,201A CURRENT FILING DATE: 2002-05-21

NUMBER OF SEQ ID NOS: 35
SOPTWARE: Patentin Ver. 2.0
SEQ ID NO 20
LENGTH: 20
TYPE: PRT
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                                                                                                                                                                                                                                                Sequence 20, Application US/09964201A Publication No. US20030091575A1 GENERAL INFORMATION:
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Matches 5; Conserv
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GENERAL INFORMATION:
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Matches 5; Conser
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APPLICANT: Van Gastel, Franciscus J.C.
ADPLICANT: Tanssen, Giselle G.
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TITLE OF INVENTION: Binding Phenol Oxidizing Enzyme-Peptide
TITLE OF INVENTION: Complexes
FILE REFERENCE: GC690-2
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CURRENT FILING DATE: 2002-09-03
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TYPE: PRT
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
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Pred. No. 1.7e+06;
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Pred. No. 1.7e+06;
1; Mismatches 0
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                                                                                                                                    09\026,276
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APPLICANT: Delon, Aprile L
APPLICANT: Lohnas, Gerald L
APPLICANT: Roberts, Steven F
APPLICANT: Roberts, Steven F
ITTLE OF INVESTION: HEAT-SHOCK FUSION-BASED VACCINE SYSTEM
FILE REFERENCE: U.S. Patent Application No. 09\026,276
CURRENT APPLICATION NUMBER: US/10/681,388
CURRENT FILING DATE: 2003-10-07
PRIOR APPLICATION NUMBER: US/09/964,201
PRIOR FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 35
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 20
LENGTH: 20
TYPE: PRT
TOCANITSM: Homo Sapiens
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US-10-681-388-20
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US-10-681-389-20
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                                                                                              ; ORGANISM: Homo sapiens
US-10-681-388-20
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Best Local Similarity 36.4
Matches 4; Conservative
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LENGTH: 20
TYPE: PRT
ORGANISM: Homo sapiens
                Query Match 32.1%;
Best Local Similarity 36.4%;
Matches 4; Conservative
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Best Local Similarity 36.4%;
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APPLICANT: Roberts, Steven F
TITLE OF INVENTION: HEAT-SHOCK FUSION-BASED VACCINE SYSTEM
FILE REFERENCE: U.S. Patent Application No. 09\026,276
CURRENT APPLICATION NUMBER: US/10/681,389
CURRENT FILING DATE: 2003-10-07
PRIOR APPLICATION NUMBER: US/09/964,201A
RRIOR FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Kenten, John H
APPLICANT: Tramontano, A
APPLICANT: Pilon, Aprile
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Kenten, John H
APPLICANT: Tramontano, A
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Pilon, Aprile L
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Pilon, Aprile L
Lohnas, Gerald L
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              Score 35; DB 4; Le Pred. No. 2.9e+02; 4; Mismatches 3;
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Pred. No. 2.9e+02;
4; Mismatches 3
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APPLICANT: Chu, Yong-Liang
APPLICANT: Chu, Yong-Liang
APPLICANT: Chu, Yong-Liang
APPLICANT: Chu, Yong-Liang
TITLE OF INVENTION: Polymeric Conjugates for Delivery of
TITLE OF INVENTION: Wid-Recognized Epitopes
TITLE OF INVENTION: Wid-Recognized Epitopes
FILE REFERENCE: 3781-001-27
CURRENT APPLICATION NUMBER: US/10/062,710
CURRENT FILING DATE: 2002-02-05
PRIOR APPLICATION NUMBER: US 60/310,498
PRIOR FILING DATE: 2001-08-08
NUMBER OF SEQ ID NOS: 232
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 50
LENGTH: 21
                                                                                                                                                                                                                            APPLICANT: DEGRECOT, ANNE S.

TITLE OF INVENTION: HIV VACCINE CANDIDATE PEPTIDES
FILE REFERENCE: 17999-001
CURRENT APPLICATION NUMBER: US/10/200,708
CURRENT FILING DATE: 2002-07-22
PRIOR APPLICATION NUMBER: US/09/351,036
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: 60/092,346
PRIOR FILING DATE: 1999-07-10
PRIOR APPLICATION NUMBER: 60/115,145
PRIOR APPLICATION NUMBER: 60/115,145
PRIOR FILING DATE: 1999-01-08
PRIOR APPLICATION NUMBER: 60/130,677
PRIOR APPLICATION NUMBER: 60/130,677
PRIOR APPLICATION NUMBER: 60/130,677
PRIOR FILING DATE: 1999-04-23
NUMBER OF SEQ ID NOS: 672
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 11
US-10-200-708-47
; Sequence 47, Application US/10200708
; Publication No. US20030180314A1
; GENERAL INFORMATION:
                                                         Query Match
Best Local Similarity
Thes 5; Conserve
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; OTHER INFORMATION: HIV Helper-T Cell Epitopes
US-10-062-710-50
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US-10-062-710-50
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Best Local Similarity
                                                                                                                                                                             ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                   TYPE: PRT
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                  LENGTH: 10
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IPHPAGLKK 9
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                                   IPHPTNIHK 10
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                                                                              1; Mismatches
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                                                                                              Score 32; DB 4;
Pred. No. 4.1e+02;
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Pred. No. 6.3e+02;
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CURRENT FILING DATE: 2002-07-22
PRIOR APPLICATION NUMBER: US/09/351,036
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: 60/092,346
PRIOR FILING DATE: 1999-07-10
PRIOR APPLICATION NUMBER: 60/115,145
PRIOR APPLICATION NUMBER: 60/130,677
PRIOR APPLICATION NUMBER: 60/130,677
                                                                                                         Query Match
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Best Local
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Publication No.
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Sequence 95, Application US/10200708
Publication No. US20030180314A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US/09/351,036
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: 60/092,346
PRIOR FILING DATE: 1998-07-10
PRIOR APPLICATION NUMBER: 60/115,145
PRIOR FILING DATE: 1999-01-08
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                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/130,677 PRIOR FILING DATE: 1999-04-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: DeGroot, Anne S.
TITLE OF INVENTION: HIV VACCINE CANDIDATE PEPTIDES
FILE REFERENCE: 17999-001
                                                                                                                                                                                                                                                  PKIOR FILING DATE: 1999-04-23
NUMBER OF SEQ ID NOS: 672
                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/200,708
CURRENT FILING DATE: 2002-07-22
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TITLE OF INVENTION: HIV VACCINE CANDIDATE PEPTIDES
FILE REFERENCE: 17999-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1999-04-23
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                                                                     Similarity 55.0 5; Conservative
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                                IPHPTNIHK 10
IPHPAGLKK 10
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                                                                                  29.4%;
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                                                                   Score 32; DB
Pred. No. 4.1e
1; Mismatches
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Pred. No. 4.1e+02;
1; Mismatches 3;
                                                               4.1e+02;
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                                                                                                    Length 10;
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RESULT 14

US-10-417-895A-36 ; Sequence 36, Application US/10417895A ; Publication No. US20040033569A1

RESULT 16

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US-10-200-708-671
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; TYPE: PRT
; ORGANISM: Human immunodeficiency virus
US-10-200-708-119
                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/200,708
CURRENT FILING DATE: 2002-07-22
PRIOR APPLICATION NUMBER: US/09/351,036
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: 60/092,346
PRIOR APPLICATION NUMBER: 60/092,346
PRIOR APPLICATION NUMBER: 60/115,145
PRIOR APPLICATION NUMBER: 60/115,145
PRIOR APPLICATION NUMBER: 60/130,677
PRIOR APPLICATION NUMBER: 60/130,677
                                                                          Best Local Similarity
Matches 5; Conserv
                                                                                                               Query Match
                                                                                                                                                                                                                                    SEQ ID NO 671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 671, Application No. US201 GENERAL INFORMATION:
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GENERAL INFORMATION:
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Best Local Similarity
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CURRENT FILING DATE: 2002-07-22
PRIOR APPLICATION NUMBER: US/09/351,036
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: 60/092,346
PRIOR FILING DATE: 1998-07-10
                                                                                                                                                                                                                                                         SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: DeGroot, Anne S.
TITLE OF INVENTION: HIV VACCINE CANDIDATE PEPTIDES
FILE REFERENCE: 17999-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/115,145
PRIOR TILING DATE: 1999-01-08
PRIOR APPLICATION NUMBER: 60/130,677
PRIOR FILING DATE: 1999-04-23
                                                                                                                                                                                                                                                                                              PRIOR FILING DATE:
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TITLE OF INVENTION: HIV VACCINE CANDIDATE PEPTIDES
                                                                                                                                                                    TYPE: PRT ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                LENGTH:
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                                   IPHPTNIHK 10
IPHPAGLKK 10
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o. US20030180314A1
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                                                                            Conservative
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                                                                                            29.4%;
                                                                          Score 32; DB 4;
Pred. No. 4.1e+02;
1; Mismatches 3
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Pred. No.
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                                                                                                             Length 10;
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US-10-808-187-2265
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                                       ; ORGANISM: Human severe acute respiratory system virus US-10-808-187-2265
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PRIOR FILING DATE: 2002-04-17
NUMBER OF SEQ ID NOS: 86
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 36
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APPLICANT: Crea, Roberto
APPLICANT: Cappuccilli,
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Best Local Similarity
                                                                                                                              SEQ ID NO 2265
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Query Match
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TITLE OF INVENTION: A DIAGNOSTIC ASSAY FOR THE HUMAN VIRUS CAUSING SEVERE ACUTE
TITLE OF INVENTION: RESPIRATORY SYNDROME (SARS)
FILE REFERENCE: V9661.0078
CURRENT APPLICATION NUMBER: US/10/808,187
CURRENT FILING DATE: 2004-03-24
CURRENT FILING DATE: 2004-03-24
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-26
PRIOR FILING DATE: 2003-03-26
PRIOR FILING DATE: 2003-03-26
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TITLE OF INVENTION: "DOPING" IN WALK-THROUG
FILE REFERENCE: 1551.2002-001
CURRENT APPLICATION NUMBER: US/10/417,895A
CURRENT FILING DATE: 2003-04-16
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                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/471,200 PRIOR FILING DATE: 2003-05-16
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NUMBER OF SEQ ID NOS: 2476
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                      ENGTH: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/459,931 FILING DATE: 2003-04-02
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/460,357 FILING DATE: 2003-04-03
                                                                                                                                                                                                                                                                                                                                        FILING DATE: 2003-04-08
APPLICATION NUMBER: 60/462,805
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                                                                                                                                                                                                                                    FILING DATE: 2003-04-23
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CHAN, KWOK HUNG
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29.4%;
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Pred. No. 5.8e
1; Mismatches
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Score 32;
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BG
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Length 14;
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US-10-103-395-109
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CURRENT FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/457,031
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-24
PRIOR TILING DATE: 2003-03-26
PRIOR APPLICATION NUMBER: 60/459,931
PRIOR APPLICATION NUMBER: 60/460,357
PRIOR APPLICATION NUMBER: 60/460,357
PRIOR APPLICATION NUMBER: 60/460,357
PRIOR APPLICATION NUMBER: 60/461,265
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                                                                                                                                                              Sequence 109, Application US/10103395 Publication No. US20020160019A1 GENERAL INFORMATION:
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SEQ ID NO 2265
LENGTH: 14
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local
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APPLICANT:
               TITLE OF INVENTION: IDENTIFICATION OF BROADLY TITLE OF INVENTION: RESTRICTED EPITOPES FILE REFERENCE; 39963-20016.01
                                                                           APPLICANT: EPIMMUNE, Inc.
APPLICANT: Sette, Alessa
APPLICANT: Sidney, John
APPLICANT: Southwood, Sc
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TITLE OF INVENTION: A HIGH-THROUGHPUT DIAGNOSTIC ASSAY FOR THE HUMAN VIRUS
TITLE OF INVENTION: CAUSING SEVERE ACUTE RESPIRATORY SYNDROME (SARS)
FILE REFERENCE: V9661.0077
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PRIOR APPLICATION NUMBER: 60/462,805
CURRENT APPLICATION NUMBER: US/10/103,395
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5; Conserv
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                                                                              Sette, Alessandro
Sidney, John
Southwood, Scott
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Pred. No. 5.
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US-10-442-909-32
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; ORGANISM: Homo sapiens
US-10-103-395-109
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PRIOR APPLICATION NUMBER: US 09/009,953
PRIOR FILING DATE: 1998-01-21
PRIOR APPLICATION NUMBER: PCT/US98/01373
PRIOR FILING DATE: 1998-01-23
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 60/036,713
PRIOR APPLICATION NUMBER: US 60/037,432
PRIOR PRIOR FILING DATE: 1997-02-07
                                                                                                                                       Sequence 32, Application US/10442909 Publication No. US20040001845A1 GENERAL INFORMATION:
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Best Local Similarity
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 19
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SEQ ID NO 109
LENGTH: 15
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Best Local
            APPLICANT: Altfeld, Marcus
APPLICANT: Yu, Xu
APPLICANT: Walker, Bruce
APPLICANT: Walker, Bruce
TITLE OF INVENTION: Cytotoxic T-Cell Epitopes of HIV-1 Virus
FILE REFERENCE: 24028-010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Chu, Yong-Liang
APPLICANT: Qiu, Jian-Tai
TITLE OF INVENTION: Polymeric Conjugates for Delivery
TITLE OF INVENTION: MHC-Recognized Epitopes
TITLE OF INVENTION: WHC-Recognized Epitopes
FILE REFERENCE: 3781-001-27
CURRENT APPLICATION NUMBER: US/10/062,710
CURRENT FILING DATE: 2002-02-05
PRIOR APPLICATION NUMBER: US 60/310,498
REIOR RELING DATE: 2001-08-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Li, Frank Q. APPLICANT: Chu, Yong-L
FILE REFERENCE: 24028-010
CURRENT APPLICATION NUMBER: US/10/442,909
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Pred. No. 6.7e+02;
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Pred. No. 6.3e+02;
1; Mismatches 3
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RESULT 23
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                                                                                                                                                                                                          GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: P
SEQ ID NO 32
                                                                                                                                                                                                                        Sequence 128, Appropriate Publication No.
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Best Local :
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Best Local :
                              PRIOR APPLICATION NUMBER: PCT/GB2004/004038
PRIOR FILING DATE: 2004-09-23
PRIOR APPLICATION NUMBER: GB 0325011.5
PRIOR FILING DATE: 2003-10-27
                                                                                                            CURRENT APPLICATION NUMBER: US/11/008,653
CURRENT FILING DATE: 2004-12-09
                                                                                                                                                APPLICANT: Schneider, Joerg
TITLE OF INVENTION: HIV Pharmaccines
FILE REFERENCE: 3742.1001-000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/285,649A
CURRENT FILING DATE: 2003-02-07
PRIOR APPLICATION NUMBER: US 08/984,099
PRIOR FILING DATE: 1997-12-03
PRIOR APPLICATION NUMBER: PCT/US96/09897
PRIOR FILING DATE: 1996-06-07
PRIOR FILING DATE: 1995-06-07
PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Stalker, David M.
APPLICANT: Pear, Julie
APPLICANT: Perez-Grau, Luis
TITLE OF INVENTION: COTTON FIBER TRANSCRIPTIONAL FACTORS
FILE REFERENCE: 15615/03/US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 18
SOFTWARE: IBM PC; Windows 2000; Microsoft Word 2000
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TYPE: PRT
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APPLICATION NUMBER: GB 0322637.0 FILING DATE: 2003-09-26
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b. US20050175627A1
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Pred. No. 7.2e+02;
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Pred. No. 8.5e+02;
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RESULT 24
US-10-430-685-95
JS-10-430-685-95
Sequence 95, Application US/10430685
Publication No. US20040039543A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: COMPUTER METHOD AND APPARATUS FOR CLASSIFYING OBJECTS
FILE REFERENCE: 63040-010210
FILE REFERENCE: 63040-010210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 95
; LENGTH: 17
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-430-685-95
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; OTHER INFORMATION: peptide from pol pool 2
US-11-008-653-128
                                                                                                                                                                                                          APPLICANT: BILLUTLA, RENUKA
APPLICANT: BRISSETTE, RENEE
APPLICANT: BRUME, ARTHUR J.
APPLICANT: SCHAFFER, LAUGE
APPLICANT: GRANDT, JAKOB
APPLICANT: GOLGSTEIN, NEIL I.
APPLICANT: SPETZLER, JANE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2003-09-24
NUMBER OF SEQ ID NOS: 174
SOPTWARE: FRBESSEQ for Windows Version 4.0
SEQ ID NO 128
LENGTH: 20
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                   Sequence 51, Application US/09962756 Publication No. US20030195147A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 55.6%;
Matches 5; Conservative
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Best Local Similarity 66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/430,685
CURRENT FILING DATE: 2003-05-06
PRIOR APPLICATION NUMBER: PCT/US01/44000
PRIOR FILING DATE: 2001-11-06
PRIOR APPLICATION NUMBER: 60/246,196
PRIOR PILING DATE: 2000-11-06
NUMBER OF SEQ ID NOS: 240
             APPLICANT: ÖSTERGAARD, SOREN
APPLICANT: HANSEN, PER HERTZ
TITLE OF INVENTION: INSULIN AND IGF-1 RECEPTOR AGONISTS AND ANTAGONISTS
FILE REFERENCE: 1878-4051US1
CURRENT APPLICATION NUMBER: US/09/962,756
CURRENT FILING DATE: 2001-09-24
PRIOR APPLICATION NUMBER: 09/538,038
PRIOR APPLICATION NUMBER: 09/538,038
PRIOR APPLICATION NUMBER: 09/146,127
FILING DATE: 1998-09-02
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GOLDSTEIN, NEIL I.
SPETZLER, JANE
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Pred. No. 8.5e+02;
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Search completed: January 26, 2006, 08:38:39 Job time: 63.75 secs
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                                                                                                          Best Local Similarity 37.5%;
Matches 6; Conservative
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SEQ ID NO 51
LENGTH: 21
                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 2227
                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: peptide
                                                                                      3 PHPTNIHKYLVCESVN 18
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PHGSNFYDWFV-EAIN
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1: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

8: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
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Copyright (c) 1993 - 2006 Compugen
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US-11-045-024-12140
US-11-045-024-13140
US-11-038-980-3
US-11-038-980-3
US-11-038-980-9
US-11-038-980-10
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 Sequence
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3294, Ap

12085, A

3376, Ap

10215, A

10215, A

10216, A

103140, A

10416, A

10516, A

10516
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US-11-045-024-1023
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13222, Ap

12094, A

12094, A

12095, Ap

12288, App

1236, App

1237, App

1238, App

1248, App

1258, App

1268, App

1278, App

1288, App

1488, App

1488, App

1428, App
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US-11-045-024-10214
Sequence 10214, Application US/11045024
Publication No. US20050271676A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                             RESULT 2
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PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-06-04
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILING DATE: 1993-08-06
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR APPLICATION NUMBER: US 08/205,713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FABESEQ for Windows Version 4.0
SEQ ID NO 3294
LENGTH: 10
Type: pom
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                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                     APPLICANT: Sette, Alessandro APPLICANT: Sidney, John APPLICANT: Southwood, Scott APPLICANT: Livingston, Briar APPLICANT: Chesnut, Robert
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TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007
CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
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                                    APPLICANT:
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APPLICATION NUMBER: US 08/347,610
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Celis,
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Chesnut, Robert
Baker, Denise Marie
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                                Baker, Denise Marie
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            Celis, Esteban
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US-11-140-417-18
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Sequence 18,
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; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS US-11-045-024-10214
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US-11-045-024-12085
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CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-06-04
PRIOR PILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-10-29
PRIOR PILING DATE: 1993-11-29
PRIOR FILING DATE: 1993-11-29
                    PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILING DATE: 1993-11-29
PRIOR FILING DATE: 1993-11-29
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Best Local
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APPLICANT:
APPLICANT:
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APPLICANT: Epimmune Inc.
TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.004
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PRIOR FILLING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
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US 08/205,713

    Mismatches

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Pred. No. 11;
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PRIOR FILING DATE: 1994-03-04 PRIOR APPLICATION NUMBER: US

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; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
US-11-045-024-3376
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PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12085
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SEQ ID NO 3376
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Best Local
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CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1993-0-05
PRIOR FILING DATE: 1993-03-05
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR PILING DATE: 1993-06-04
PRIOR PILING DATE: 1993-08-06
PRIOR PILING DATE: 1993-08-06
PRIOR PILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-08-06
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IPHPAGLKK 10
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Southwood, Scott
                                                                                   Conservative
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Pred. No.
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Pred. No. 13;
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PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FASTSEQ for Windows Version 4.
SEQ ID NO 10191
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Best Local Similarity
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APPLICANT:
TITLE OF INVENTION: Inducing Cellular Responses TITLE OF INVENTION: Virus-1 Using Peptide and PEILE REFERENCE: 2060.0040007 CURRENT APPLICATION NUMBER: US/11/045,024 CURRENT PILING DATE: 2005-01-28
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CURRENT FILING DATE: 2005-01-28
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FILE REFERENCE: 2060.0040007
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FILING DATE: 1993-03-05
APPLICATION NUMBER: US 08/073,205
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FILING DATE: 1993-11-29
APPLICATION NUMBER: US 08/159,339
FILING DATE: 1993-11-29
APPLICATION NUMBER: US 08/205,713
FILING DATE: 1994-03-04
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APPLICATION NUMBER: US 08/103,396
FILING DATE: 1993-08-06
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                                                                                                             Kubo, Ralph
Grey, Howard M...
                                                                                             Epimmune Inc.
                                                                                                                                                Celis, Esteban
                                                                                                                                                                                     Livingston, Bri
Chesnut, Robert
                                                                                                                                                                                                                            Southwood, Scott
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                                                                                                                                                                Baker, Denise Marie
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Celis, Esteban
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Pred. No. 13;
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Nucleic A
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US-11-045-024-12063
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PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 10215
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Publication No. US20050271676A1
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 12063
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PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-06-04
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILING DATE: 1993-08-06
PRIOR FILING DATE: 1993-08-06
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APPLICANT:
APPLICANT:
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                                       PRIOR FILING DATE: 1994-12-01
NUMBER OF SEQ ID NOS: 14528
                                                                              PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
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PRIOR FILING DATE: 1993-03-05
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APPLICATION NUMBER: US 08/103,396
FILING DATE: 1993-08-06
APPLICATION NUMBER: US 08/159,184
FILING DATE: 1993-11-29
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FILING DATE: 1999-10-05
APPLICATION NUMBER: US 08/027,146
                                                                                                                                         FILING DATE: 1993-11-29
APPLICATION NUMBER: US 08/159,339
FILING DATE: 1993-11-29
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APPLICATION NUMBER: US 08/073,205
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Grey, Howard M.
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Pred. No.
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US-11-045-024-13140

Sequence 13140, Application US/11045024
Publication No. US20050271676A1
GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John

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TITLE OF INVENTION: Unducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007
CCURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR FILING DATE: 1993-08-04
PRIOR FILING DATE: 1993-06-04
PRIOR FILING DATE: 1993-06-04
PRIOR FILING DATE: 1993-06-06
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR PILING DATE: 1993-11-29
PRIOR PILING DATE: 1993-11-29
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Best Local Similarity
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12086
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PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
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Local Similarity 55.6%;
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  IPHPAGLKK 11
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Celis,
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                                                                                             Score 32; DB
Pred. No. 13;
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SOFTWARE: FABTSEQ
SEQ ID NO 13140
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                                                                SEQ ID NO 3
                                                                                                               APPLICANT: Michel Perricaudet
TITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
TILE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
TILE REFERENCE: p26,992-C USA:
CURRENT FILING DATE: 2005-01-20
PRIOR APPLICATION NUMBER: US 09/791,524
PRIOR FILING DATE: 2001-02-22
PRIOR FILING DATE: 1909-08-27
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/098,028
PRIOR FILING DATE: 1998-08-27
PRIOR FILING DATE: 1998-08-27
PRIOR FILING DATE: 1998-08-27
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APPLICANT: Emmanuelle Vigne
                                                                                                           NUMBER OF SEQ ID NOS:
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PRIOR FILING DATE: 1993-11-29
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PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
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CURRENT FILING DATE: 2005-01-28
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ORGANISM: Adenovirus
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FastSEQ for Windows Version 4.0
                                                                                  PatentIn version 3.2
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Patrice Yeh
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Pred. No.
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Matches 5; Conserv
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Best Local
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Publication No.
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                               PRIOR APPLICATION NUMBER: US 09/791,524
PRIOR FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: PCT/IB99/01524
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/098,028
PRIOR FILING DATE: 1998-08-27
                                                                                                                                   APPLICANT: Patrice Yeh
APPLICANT: Michel Perricaudet
TITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
FILE REFERENCE: P36,992-C USA
CURRENT APPLICATION NUMBER: US/11/038,980
CURRENT FILING DATE: 2005-01-20
          PRIOR FILING DATE: 1998-08-27
NUMBER OF SEQ ID NOS: 165
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PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/098,028
PRIOR FILING DATE: 1998-08-27
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TYPE: PRT
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WARE: PatentIn version 3.2
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No. US20060002893A1
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Jean-Francois De
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Patrice Yeh
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Pred. No. 29;
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CURRENT APPLICATION NUMBER: US/11/038,980
CURRENT FILING DATE: 2005-01-20
PRIOR APPLICATION NUMBER: US 09/791,524
PRIOR FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: PCT/IB99/01524
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/098,028
PRIOR APPLICATION NUMBER: US 60/098,028
PRIOR FILING DATE: 1998-08-27
NUMBER OF SEQ ID NOS: 165
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 10
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US-10-859-643-123
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APPLICANT: Challita-Eid, Pia M.
APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
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                                                                                                           APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Entitled 161P2F10B Useful in Treatment
TITLE OF INVENTION: Cancer
FILE REFERENCE: 511582006203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Jean-Francois Dedieu
APPLICANT: Martine Latta
APPLICANT: Patrice Yeh
APPLICANT: Patrice Yeh
APPLICANT: Michel Perricaudet
TITLE OF INVENTION: Targeted Adenovirus Vectors For Delivery Of Heterologous Genes
FILE REFERENCE: P26,992-C USA
CURRENT APPLICATION NUMBER: US/10/859,643
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 765
SOFTWARE: FASTSEQ for Windows Version 4.0
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TYPE: PRT
ORGANISM: Adenovirus
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No. US20060002893A1
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Pred. No. 34;
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RESULT 16
US-11-097-912-123
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APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
APPLICANT: Hubert, Rene S.
APPLICANT: Morrison, Karen Jane
APPLICANT: Jakobovits, Aya
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PRIOR ETLING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 765
SOFTWARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 123
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Best Local Similarity 75.0
Matches 6; Conservative
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APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Jakobovits, Aya
APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED 161P2F10B
TITLE OF INVENTION: USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 511582006204
CURRENT APPLICATION NUMBER: U$(11/097,912
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR APPLICATION NUMBER: US 10/05,480
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 765
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APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: NUCLERIC ACID AND CORRESPONDING PROTEIN NAMED 161P2F10B
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 511582006205
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CURRENT FILING DATE: 2005-04-01
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ORGANISM: Homo Sapien
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Morrison, Karen Jane Meyrick
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75.0%;
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Pred. No. 5.9e+04;
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Pred. No. 5.9e+04;
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; ORGANISM: Homo Sapien
US-10-859-643-164
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; ORGANISM: Homo Sapien
US-11-097-912-123
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 164
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Matches 6; Conserv
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                                                          APPLICANT: Hubert, Rene S.
APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Morrison, Karen Jane Meyrick
APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 161P2F10B
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 511582005205
CURRENT APPLICATION NUMBER: US/11/097,864
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 10/062,109
PRIOR FILING DATE: 2002-01-31
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APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
TITLE OF INVENTION: Entitled 161P2F10B Useful in Treatment and Detection of
TITLE OF INVENTION: Cancer
TITLE OF INVENTION: Cancer
PRIOR APPLICATION NUMBER: US 10/005,480 PRIOR FILING DATE: 2001-11-07 NUMBER OF SEQ ID NOS: 765
                                                                                                                                                                                                                                                                                               APPLICANT: Challita-Eid, Pia M. APPLICANT: Raitano, Arthur B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/859,643
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: US 10/005,480
PRIOR FILING DATE: 2001-11-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version
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lication No. US20060002993A1
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Faris, Mary
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Faris, Mary
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75.0%;
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Pred. No. 29
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Pred. No. 5.9e+04;
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; ORGANISM: Homo Sapien
US-11-097-912-164
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 164
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Best Local Similarity
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Best Local :
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CURRENT APPLICATION NUMBER: US/10/966,483
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: US 60/511,919
PRIOR FILING DATE: 2003-10-15
PRIOR APPLICATION NUMBER: US 60/511,719
PRIOR APPLICATION NUMBER: US 60/511,719
PRIOR APPLICATION NUMBER: US 60/532,666
PRIOR FILING DATE: 2003-12-24
                                                                                                                                                                                                    APPLICANT: Kinch, Michael S.
APPLICANT: Kiener, Peter A.
APPLICANT: Bruckheimer, Elizabeth
APPLICANT: Dubensky, Jr. Thomas W.
APPLICANT: Cook, David N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED 161P2F10B
TITLE OF INVENTION: USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 511582006204
CURRENT APPLICATION NUMBER: US/11/097,912
CURRENT FILING DATE: 2005-04-01
                                                                                                                                                                FILE REFERENCE: 10271-146
                                                                                                                                                                                   TITLE OF INVENTION: LISTERIA-BASED EphA2 VACCINES
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TYPE: PRT
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TYPE: PRT
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Faris, Mary
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75.0%;
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Pred. No. 2
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Pred. No. 29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-11-166-412-88
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LENGTH: 14
TYPE: PRT
               GENERAL INFORMATION:
APPLICANT: Sahin Dr., Ugur
APPLICANT: Tureci Dr., Ozlem
APPLICANT: KOSlowski Dr., Michael
APPLICANT: KOSLowski Dr., Michael
APPLICANT: KOSLOwski Dr., Michael
TITLE OF INVENTION: Genetic Products Differentially Expressed in Tumors and Use There
FILE REFERENCE: 342-3PCT
CURRENT APPLICATION NUMBER: US/10/506,443A
CURRENT APPLICATION 100
NUMBER OF SEQ ID NOS: 100
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SEQ ID NO 17
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Publication No.
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Best Local Similarity
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Best Local (
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APPLICANT: Tomme, Peter H. M.
APPLICANT: Tomme, Peter H. M.
APPLICANT: Tomme, Peter H. M.
TITLE OPT INVENTION: Methods and Compositions To Promote Bone Homeostasis
FILE REFERENCE: P27,927-D USA
CURRENT APPLICATION NUMBER: US/11/166,412
CURRENT FILING DATE: 2005-06-24
PRIOR APPLICATION NUMBER: 60/582,704
PRIOR APPLICATION NUMBER: 60/630,449
PRIOR APPLICATION NUMBER: 60/630,449
PRIOR APPLICATION NUMBER: 60/673,206
PRIOR APPLICATION NUMBER: 60/673,206
PRIOR FILING DATE: 2005-04-20
NUMBER OF SEQ ID NOS: 231
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PRIOR FILING DATE: 2004-10-01
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 2004-10-07
NUMBER OF SEQ ID NOS: 72
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PatentIn version 3.1
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No. US20060014231A1
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nilarity 41.7%;
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Pred. No. 50;
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; SEQ ID NO 18
; LENGTH: 17
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-506-443A-18
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US-11-045-024-1627
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CURRENT APPLICATION NUMBER: US/10/723,207
CURRENT FILING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: 09/701,623
PRIOR FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: PCT/US99/13959
PRIOR FILING DATE: 1999-06-21
PRIOR APPLICATION NUMBER: 09/100,287
PRIOR APPLICATION NUMBER: 09/100,287
PRIOR APPLICATION NUMBER: 09/100,287
PRIOR FILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
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Best Local Similarity 50.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                        APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Livingston, Briar
                TITLE OF INVENTION: Inducing Cellular Responses to Human Immunodeficiency TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions FILE REFERENCE: 2060.0040007
                                                                                             APPLICANT:
                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF TITLE OF INVENTION: ALLERGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Wang, Chang Yi APPLICANT: Walfield, Alan M.
                                                                             APPLICANT:
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CURRENT APPLICATION NUMBER: US/11/045,024
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                                                                           Epimmune Inc.
                                                                                             Kubo, Ralph
Grey, Howard M.
                                                                                                                               Baker, Denise Marie
Celis, Esteban
                                                                                                                                                                          Chesnut, Robert
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Pred. No. 60;
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TITLE OF INVENTION: Unducing Cellular Responses to Human Immunodeficiency
TITLE OF INVENTION: Virus-1 Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0040007
CURRENT APPLICATION NUMBER: US/11/045,024
CURRENT FILING DATE: 2005-01-28
PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/103,396
PRIOR FILING DATE: 1993-06-04
PRIOR FILING DATE: 1993-01-05
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/159,184
PRIOR APPLICATION NUMBER: US 08/159,184
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TYPE; PRT;
; ORGANISM; HUMAN IMMUNODEFICIENCY VIRUS
US-11-045-024-1627
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SEQ ID NO 1627
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PRIOR APPLICATION NUMBER: US 08/159,339
PRIOR FILING DATE: 1993-11-29
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Livingston, Brian
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PRIOR FILING DATE: 1994-03-04
PRIOR APPLICATION NUMBER: US 08/347,610
PRIOR FILING DATE: 1994-12-01
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR APPLICATION NUMBER: US 09/412,863
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 08/027,146
PRIOR FILING DATE: 1993-03-05
PRIOR APPLICATION NUMBER: US 08/073,205
PRIOR APPLICATION NUMBER: US 08/073,205
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PRIOR FILING DATE: 1993-11-29
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4; Conserv
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; LENGTH: 8
; TYPE: PRT
; ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
US-11-045-024-4340

Query Match
Best Local Similarity 57.1%; Score 28; DB 7; Length 8;
Best Local Similarity 57.1%; Pred. No. 5.9e+04;
Matches 4; Conservative 1; Mismatches 2; Indels 0; Gaps
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A56786
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Ig H chain V-D-J r
hypothetical TEL/M
trypsin (EC 3.4.21
beta 3-adrenergic
glycogen(starch) s calmodulin, vasoac GTP-binding protei entactin/nidogen - chondroitin sulfat glandular kallikre interferon alpha r cystatin CI-4a - m pimeloyl-CoA synth T-cell receptor be T-cell receptor be RNA-directed RNA p venombin B (EC 3.4 iodothronine 5'-mo 15K protein B - ra phosphatidylinosit alpha-conotoxin Cn thyroglobulin - pi thyroglobulin - pi thyroglobulin - sh trehalase inhibito phospheenolpyruvat cytochrome-c oxida Ig heavy chain CRD cell surface glyco hemocyanin chain 3 protein QA300050 gluten - wheat photosystem II ch. fimbrial antigen intracrystalline T-cell receptor k
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T-cell receptor be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Accession: A60929
R;Kohler-Staub, D.; Hartmans, S.; Gaelli, R.; Suter, F.; Leisinger, J. Gen. Microbiol. 132, 2837-2843, 1986
A;Title: Evidence for identical dichloromethane dehalogenases in di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              dichloromethane dehalogenase (EC 4.5.1.3) - Hyphomicrobium sp. (ATCC 43129) (fragment)
                                                                                                           A; Molecule type: DNA
A; Residues: 1-11 < GUZ>
                                                                                                                                                                                                                                                           celf protein - Escherichia coli (fragment)
C;Species: Escherichia coli
C;Date: 25-Dec-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
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C;Keywords: carbon-halide lyase
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J. Gen. Microbiol. 132, 2837-2843, 1986
A;Title: Evidence for identical dichloromethane dehalogenases in different methylotrophi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Pseudomonas sp.
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text
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                                                        A;Gene:
                                                                                         A; Cross-references: UNIPARC: UPI000017AA33
                                                                                                                                              A;Reference number: S42587;
A;Accession: S42587
                                                                                                                                                                                                                          R;Guzzo, A.; DuBow, M.S
                                                                                                                                                                                                                                           C; Accession: S42587
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Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004;

Accession: A60929
                                                                                                                                                                                     Title: A luxAB transcriptional
                                                                            Genetics
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Best Local Similarity
                 Query Match
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A luxAB transcriptional fusion to
ce number: S42587; MUID:94166755;
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85.7%;
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Pred.
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C;Accession: H64711

R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R. Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKen son, J.D.; Kelley, J.M.; Cotton, M.D.; Weldman, J.M.; Fujii, C.; Bowman, C.; Watthey, Nature 388, 539-547, 1997

A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori. A;Reference number: A64520; MUID:97394467; PMID:9252185

A;Accession: H64711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: protein
A;Residues: 1-5;6-10;11-18 <INO>
A;Residues: 1-5;6-10;11-18 <INO>
A;Cross-references: UNIPARC:UPI000017B3BC; UNIPARC:UPI000017B3BD; UNIPARC:UPI000017B3BE
A;Experimental source: strain var. macrosporus
C;Keywords: aspartic proteinase; hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A,Title: Heterologous expression and site-directed mutagenesis studies on A,Reference number: S65454; MUID:96235238; PMID:8647118 A,Accession: S65454
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R;Inoue, H.; Hayashi,
Eur. J. Biochem. 237,
R;Cooke, R.J.; Bjoernestedt, R.; Douglas, K.T.; McKie, J.H.; King, M.D.; Coles, B.; Ket Biochem. J. 302, 383-390, 1994
A;Title: Photoaffinity labelling of the active site of the rat glutathione transferases A;Reference number: S55307; MUID:94379965; PMID:8092989
                                                                                          glutathione transferase 3-3 - rat (fragments)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 07-May-1999
C;Accession: S55307
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C;Date: 28-Oct_1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
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;Date: 09-Aug-1997 #sequence_revision 09-Aug-1997
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                                                                      Bjoernestedt, R.; Douglas, K.T.; McKie, J.H.; King, M.D.; Coles,
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Pred. No. 1e+03;
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Pred. No. 7.2e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypothetical protein At2g31090 [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004 C;Accession: D84716
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A; Residues: 1-20 <STO>
                                                                                                                                                                                                                                                                                                                         A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: D84716
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A;Molecule type: protein
A;Residues: 1-16 <COO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references: UNIPARC: UPI000017C91E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Davison, M.D.; Rixon, F.J.; Davison, A.J.
Gen. Virol. 73, 2709-2713, 1992

Title: Identification of genes encoding two capsid proteins
Reference number: PQ0544; MUID:93019027; PMID:1328483
Accession: PQ0548
                                                                                                                                                                                             Genetic:
Gene: At2g31090
Man mosition: 2
                                                                                                                                                                                                                                                                                                      Status: preliminary
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;Experimental source: strain 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Molecule type: protein ;Residues: 1-19 <DAV>
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Best Local
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Date: 03-Feb-1994 #sequence_revision
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                                                                                     12 LYCESVNG
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                                                    LVCDCTSG 14
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                                                                                                                       Score 25; DB
Pred. No. 1.7e
2; Mismatches
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Pred. No.
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Pred. No. 1.1e+03;
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                                                                                                                 DB 2;
1.7e+03;
2;
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R;Bergman, T.; Agerberth, B.; CFEBS Lett. 283, 100-103, 1991
A;Title: Direct analysis of per A;Reference number: S15778; MUI A;Accession: S15778
A;Status: preliminary
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N;Alternate names: blastogen A; scarlet fever toxin
C;Species: Streptococcus pyogenes
C;Date: 10-Nov-1992 #sequence_revision 10-Nov-1992.#text_change 31-Dec-2004
                                                             A;Molecule type: protein A;Residues: 9-17 <FEB2>
                                                                                                                        A; Cross-references: A; Accession: S15779
                                                                                                                                                                 A;Molecule type: protein
A;Residues: 1-8 <FEB1>
                                                                                                                                                                                                                                                                                                                                                   C;Species: Bos primigenius indicus x Bos primigenius taus
C;Date: 19-Mar-1997 #sequence_revision 16-Jul-1999 #text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biocnemistry (N.Y.) 53, 183-190, A;Title: Low-molecular-weight pe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Tuichibaev, M.U.; Akhmedova, Biochemistry (N.Y.) 53, 183-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   venom protein HR-3 - oriental hornet (fragment)
C;Species: Vespa orientalis (oriental hornet)
C;Species: 29-Jan-1993 #sequence_revision 29-Jan-1993 #text_change 09-Jul-2004
C;Date: 29-Jan-1993 #sequence_revision 29-Jan-1993 #text_change 09-Jul-2004
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C;Superfamily: enterotoxin B
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Infect. Immun. 57, 1865-1867, 1989
A;Tille: Group A streptococcal pyrogenic exotoxin (scarlet A;Reference number: A60108; MUID:89254013; PMID:2498210
                                                                                                  A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                 insulin chain B - bovine (fragments)
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                                         Cross-references: UNIPARC:UPI000015BC0B
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                        Superfamily:
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Best Local
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1-9 <TUI>
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pancreas
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Pred. No.
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No. 2.8e+05;
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2.8e+05;
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N;Alternate names: 14K yolk protein
C;Species: Theromyzon tessulatum (duck leech)
C;Date: 19-Mar_1997 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
                                                                                                                        RESULT 14
I50153
                                                                                                                                                                                                                                                                                                                                                                                    R:Tsugita, A.; Kamo, M.; Kawakami, M.; Ohki, Y. submitted to JIPID, December 1995
A;Description: Two dimensional electrophoresis of A;Reference number: PN0173
A;Accession: PN0175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                glutathione transferase (EC 2.5.1.18) 1 - Arabidopsis thaliana (fragment) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 22-Dec-1995 #sequence_revision 08-Feb-1996 #text_change 23-Mar-2001 C;Accession: PN0175
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PN0175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Baert, J.L.; Britel, M.; Sautiere, P.; Malecha, J. Eur. J. Biochem. 209, 563-569, 1992
A;Title: Ovohemerythrin, a major 14-kDa yolk protein distinct A;Reference number: S29264; MUID:93049299; PMID:1425663
A;Accession: S29264
Mol. Cell. Biol. 4,
A;Title: Isolation a
A;Reference number:
                                                         alpha-actin - chicken (fragment)
C;Species: Gallus (chicken)
C;Date: 13-Sep-1996 #sequence_revision
C;Accession: I50153
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                                            R;Chang,
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A;Residues: 1-18 <TSU>
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                                                        Accession:
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, 2498-2508, 1984 and characterization of : I50153; MUID:85085956;
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100.0%;
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Pred. No. 2.1e-
1; Mismatches
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Pred. No.
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                                                                        13-Sep-1996 #text_change
six different
PMID:6513927
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2e+03;
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2.1e+03
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                chicken
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 Ig H chain
C;Species:
                                PH1604
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A;Note: sequence extracted from NCBI backbone (NCBIP:107297)
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase; venom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: A45655
R;Chiou, S.H.; Hung, C.C.; Lin, C.W.
B;Chiou, S.H.; Hung, C.C.; Lin, C.W.
Biochem. Int. 56 105-112, 1992
A;Title: Isolation of a crotalase-like protease with alpha-fibrinogenase activity from A;Reference number: A45655, MUID:92313365; PMID:1616487
A;Accession: A45655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-19 <CHA>
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S38736
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A45655
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                                                                                                                                                                                                                                                                     FEBS Lett. 332, 243-246, 1993
A;Title: Purification and antipathogenic activity of lipid A;Reference number: S38736; MUID:94009709; PMID:8405465
                                                                                                                                                                                                                                                                                                                                                                             lipid transfer protein al - Arabidopsis thaliana (fragment) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 19-May-1994 #sequence_revision 19-Jan-1996 #text_ch
                                                                                                                                                                                                         A;Residues: 1-13 <SEG>
A;Cross-references: UNIPARC:UPI000017AFCF
                                                                                                                                                                                                                                                                                                                           R;Segura, A.; Moreno, M.;
FEBS Lett. 332, 243-246, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q9PS55; UNIPARC:UPI00000FB75C A;Experimental source: venom
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RESULT 17
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A;Residues: 1-20 <CHI>
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Pred. No. 2.4e
2; Mismatches
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Pred. No. 2.
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Pred. No. 2.3e+03;
3; Mismatches 1
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V-D-J region Mus musculus

(wild-type clone 327)
(house mouse)

mouse

(fragment)

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R;Pearson, L.; Meagher, R.B.
Plant Mol. Biol. 14, 513-526, 1990
A;Title: Diverse soybean actin transcripts contain a large
A;Reference number: S15754; MUID:91346640; PMID:2102831
A;Accession: S15754.
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S15754
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C;Accession: PH1604
R;Levinson, D.A.; Campos-Torres, J.; Leder,
J. Exp. Med. 178, 317-329, 1993
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                                                                                                            R;Meyer, M.; Granderath, K.; Andreesen, Eur. J. Blochem. 234, 184-191, 1995
                                                                                                                                                                                                                                                                                                                              83K protein -
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A; Residues: 1-17 < PEA>
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C;Datc: 04-Dec-1992 #sequence revision 04-Dec-1992 #text_change 05-Oct-2004
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A; Residues: 1-16 <LEV>
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Matches
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Best Local
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                                                              Matches
                                                                                                                                                                                                                                   Title: Purification and
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Similarity 44.4%;
4; Conservative
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Pred. No. 2.9e+03;
2; Mismatches 2;
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Pred. No. 2.9e+03;
3; Mismatches 0
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Pred. No. 2.7e+03;
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RESULT 21
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A58589
                                                               A;Reference number: A37196;
A;Accession: B37196
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A;Molecule type: prote
A;Residues: 1-10 <ALT>
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A;Note: sequence confirmed by chemical synthesis
C;Comment: This alpha-conotoxin, as an acetylcholine receptor inhibitor,
C;Superfamily: alpha-conotoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Martinez, J.S.; Olivera, B.M.; Gray, W.R.; Craig, A.G.; Groebe, D.K.; Abramson, S. Biochemistry 34, 14519-14526, 1995
A;Title: alpha-Conotoxin EI, a new nicotinic acetylcholine receptor antagonist with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              alpha-conotoxin EI - cone shell (Conus ermineus)
C;Species: Conus ermineus (ermine cone)
C;Date: 10-Sep_1999 #sequence_revision 10-Sep-1999 #text_change 09-Jul-2004
C;Accession: A58589
                                                                                     A;Title: Primary structure and biological activity of bradykinin A;Reference number: A37196; MUID:90351557; PMID:2386615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Title: Characterisation of peptide-N4-(N-acetyl-beta-glucosaminyl)asparagine
A;Reference number: A59272; MUID:98181894; PMID:9523720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eur. J. Biochem. 252, 118-123, 1998
A;Title: Characterisation of peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                F;4-10,5-18/Disulfide bonds: #Status experimental F;18/Modified site: amidated carboxyl end (Cys) #status experimental
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                     ;Status: preliminary
;Molecule type: proto
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Best Local
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9, 221-227,
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Pred. No. 3.7e+03;
2; Mismatches 4
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RESULT 23
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C;Keywords: pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
mannose-1-phosphate guanylyltransferase (EC 2.7.7.13) 43K alpha chain - N;Alternate names: GDP-mannose pyrophosphorylase 43K alpha chain C;Species: Sus scrofa domestica (domestic pig)
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A;Title: Six novel tachykinin- and bombesin-related peptides from the skin of the Austra A;Reference number: A60409; MUID:90287814; PMID:2356157

A;Accession: B60409
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C;Keywords: amidated carboxyl end; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;1/Modified site: amidated carboxyl end (Met) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: protein A; Residues: 1-11 <SIM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Reference number: A60409; A; Accession: D60409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Title: Six novel tachykinin- and bombesin-related peptides from the skin of the Austra A;Reference number: A60409; MUID:90287814; PMID:2356157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Accession: D60409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kassinin-like peptide K-III - frog (Pseudophryne guentheri)
C;Species: Pseudophryne guentheri
C;Date: 30-Jan-1993 #sequence_revision 30-Jan-1993 #text_change 16-Aug-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         kassinin-like peptide K-I - frog (Pseudophryne guentheri)
C;Species: Pseudophryne guentheri
C;Bate: 30-Jan-1993 #sequence_revision 30-Jan-1993 #text_change 16-Aug-2004
C;Accession: B60409
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C;Date: 25-Feb-1994 #sequence_revision 12-Aug-1996 #text_change 12-Aug-1996 C;Accession: B47415 R;Szumilo, T.; Drake, R.R.; York, J.L.; Elbein, A.D. J. Biol. (Chem. 268, 17943-17950, 1993 A;Title: GDP-mannose pyrophosphorylase. Purification to homogeneity, propert A;Reference number: A47415; MUID:93352609; PMID:7688733 A;Accession: B47415 A;Molecule type: protein A;Reference pyrotein A;Residues: 1-13 <SZU-A;Recession: B47415 ASU-A;Residues: 1-13 <SZU-A;Recession: B47415 ASU-A;Recession: B47415 ASU-A;R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Description: generates GDP-mannose and pyrophosphate from mannose-1-phosphate and GTP A;Note: also catalyzes synthesis of GDP-glucose from glucose-1-phosphate (EC 2.7.7.34 ac C;Keywords: blocked amino end; nucleotidyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPARC:UPI000017C470
A;Experimental source: liver
C;Complex: The enzyme appears to be a hetc
C;Function:
Search completed: January 26, 2006, 08:05:05
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Kohler-Staub D., Hartmans S., Gaelli R.,
Evidence for identical dichloromethane
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microbiol. 132:2837-2843(1986).
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Kohler-Staub D., Hartmans S., Gaelli R.,
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DE Gag protein (Fragment).
QS Human T-lymphotropic virus
QC Viruses; Retroid viruses; R
QX NCBI TAXID=11908;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RX MEDLINE=87111460; PubMed=30
RA Hiramatsu K., Nishida J., N
RT "Molecular cloning of the c
RT leukaemia virus type I: A n
RT region.";
RL J. Gen. Virol. 68:213-218(1)
DR EMBL; X04800; CAA28492.1;
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Human T-lymphotropic virus
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MEDLINE=87111460; PubMed=3027244;
Hiramatsu K., Nishida J., Naito A., Yoshika
"Molecular cloning of the closed cirular pu
leukaemia virus type I: A new open reading
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Viruses; Retroid viruses; Retroviridae; Deltaretrovirus.
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MEDLINE=85211032; PubMed=2987530;
Ratner L., Josephs S.F., Starcich
Gallo R.C., Wong-Staal F.;
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Hypothetical protein.
Schistosoma japonicum (Blood fluke).
Schistosomatoidea; Platyhelminthes; T.
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Hypothetical protein.
SEQUENCE 19 AA; 2259 MW:
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SEQUENCE
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Eukaryota; Fungi; Ascomycota; Pe
Sordariomycetidae; Sordariales;
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EMBL; M10977; AAA46203.1; -; Genomic_DNA.
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Homo.
NCBI_TaxID=9606;
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                                               Homo sapiens (Human)
                                                                                  01-NOV-1996 (TrEMBLrel. 01, 01-NOV-1996 (TrEMBLrel. 01, 01-MAY-1999 (TrEMBLrel. 10, STS protein (Fragment).
                                                                                                                                       Q16244 HUMAN
Q16244;
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transcriptomic, and proteomic analyses.";
Science 307:82-86(2005).
-i- CAUTION: The sequence shown here is derived !
-i- CAUTION: The sequence shown here shotgun (WGS)
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Hall N., Karras M., Raine J.D., Carl
Berriman M., Florens L., Janssen C.S
James K., Rutherford K., Harris B.,
James K., Rutherford M., Doggett J., T
Quail M.A., Ormond D., Doggett J., T
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13-SEP-2005 (TrEMBLrel. 31, Last sequence up
13-SEP-2005 (TrEMBLrel. 31, Last annotation
13-SEP-2005 (TrEMBLrel. 31, Last annotation
Hypothetical protein (Fragment).
ORFNames=PC400586.00.0;
ORFNames=PC400586.00.0;
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NON TER 10 1
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                                      Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=5825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Alveolata;
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EMBL; AABX01000220; EAA32832.1;
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NUCLEOTIDE SEQUENCE.
NUCLEOTIDE SEQUENCE.
MEDLINE=95038775; PubMed=7951263;
Yen P.H., Ferrero G.B., Chinault A.C
"Characterization of the deletion br
steroid sulfatase deficiency.";
Hum. Mutat. 4:76-78(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      p41495;
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
05-JUL-2004 (Rel. 44, Last annotation update)
Trypsin-modulating oostatic factor (TMOF).
Sarcophaga bullata (Grey flesh fly) (Neobellieria bullata).
Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota;
Neoptera, Endopterygota, Diptera, Brachycera, Muscomorpha, Oestroidea,
Sarcophagidae, Sarcophaga, Neobellieria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SARBU
TMOF_S
                                                                                                                                                                                                                                                                                                                                                                                                             (Sarcophaga) bullata.";
Regul. Pept. 50:61-72(1994)
-!- FUNCTION: Has an oostat
                                                                                                                                                                                                              Direct protein sequencing; SEQUENCE 6 AA; 695 MW;
                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE-94211930; PubMed-8159807; DOI=10.1016/0167-0115(94)90192-9; Bylemans D., Borovsky D., Hunt D.F., Shabanowitz J., Grauwels L.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=7385;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequencing and characterization of trypsin modulating oostatic (TMOF) from the ovaries of the grey fleshfly, Neobellieria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Loof A.;
                                                                                                                                                                                                                                                                                                                                                   development.

DEVELOPMENTAL STAGE: Synthesized and released from follicular
                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: Has an oostatic activity. Inhibits trypsin biosynthesis in the midgut which indirectly reduces the vitellogenin concentration in the hemolymph resulting in inhibition of oocyte
                                                                                                                                                                                                                                                                                                                                         epithelium after a blood meal.
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  (TrEMBLrel. 30,
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Berriman M., Florens L., Janssen C.S., Pain A., Christophides G.K.
James K., Rutherford K., Harris B., Harris D., Churcher C.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.
Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.;
"A comprehensive survey of the Plasmodium life cycle by genomic,
transcriptomic, and proteomic analyses.";
Science 307:82-86(2005).
Science 307:82-86(2005).
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                                                                                                                                                                                                Hypothetical protein.
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Eukaryota; Alveolata;
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Q4YCP3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Streptocarpus venosus.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Gesneriaceae; Cyrtandroideae; Didymocarpeae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=5821;
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ORFNames=PB100163.00.0;
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Plant Cell 17:430-443(2005).
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Harrison J., Moller M., Langdale J., Cronk G
"The Role of KNOX Genes in the Evolution of
                                                                                                                                                                            SEQUENCE
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; CAAI01006597; CAI04242.1;
                                                                                                                                                                                                                                                                                   EMBL/GenBank/DDBJ whole genome shotgun
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Morphological
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Plant Cell 17:430-443(2005).
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Harrison J., Moller M., Langdale J., Cronk Q.,
"The Role of KNOX Genes in the Evolution of Mo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Streptocarpus glandulosissimus.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; aster
lamiids; Lamiales; Gesneriaceae; Cyrtandroideae; Didymocarpeae;
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PubMed=15659624; DOI=10.1105/tpc.104.028936;
Harrison J., Moller M., Langdale J., Cronk Q
"The Role of KNOX Genes in the Evolution of P
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Epermatophyta; Magnollophyta; eudicotyledons; core eudicots; aster. lamiids; Lamiales; Gesneriaceae; Cyrtandroideae; Didymocarpeae;
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10-MAY-2005 (TrEMBLrel. 30,
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Plant Cell 17:430-443(2005).
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Plant Cell 17:430-443(2005).
EMBL; AY662116; AAW33770.1; -;
InterPro; IPR001356; Homeobox.
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Plant Cell 17:430-443(2005).
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Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.;
"The Role of KNOX Genes in the Evolution of Morphological
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Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.;
"The Role of KNOX Genes in the Evolution of Morphological Novelty
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Q5EX91,9LAMI PRELIMINARY;
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ProDom; PD000010; Homeobox;
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EMBL; AY662112; AAW33766.1;
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Plant Cell 17:4
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PubMed=15659624; DOI=10.1105/tpc.104.028936;
Harrison J., Woller M., Langdale J., Cronk Q
"The Role of KNOX Genes in the Evolution of I
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; Spermatophyta; Magnoliophyta; eudicotyledons; Didymocarpeae; lamiids; Lamiales; Gesneriaceae; Cyrtandroideae; Didymocarpeae;
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EMBL; AY662113; AAW33767.1; -;
InterPro; IPR001356; Homeobox.
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Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.;
"The Role of KNOX Genes in the Evolution of Morphological Novelty
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
Lamiids; Lamiales; Gesneriaceae; Cyrtandroideae; Didymocarpeae;
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PubMed=15659624; DOI=10.1105/tpc.104.028936;
Harrison J., Moller M., Langdale J., Cronk Q
"The Role of KNOX Genes in the Evolution of Streptocarpus.";
Plant Cell 17:440-443(2005).
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Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.;
"The Role of KNOX Genes in the Evolution of Morphological Novelty in
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NCBI_TaxID=167276;
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Eukaryota; Viridiplantae;
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Plant Cell 17:430-443(2005).
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Plant Cell 17:430-443(2005).
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STM2 protein (STM1 protein) (Fragment).
Streptocarpus burundianus.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiales; Lamiales; Gesneriaceae; Cyrtandroideae; Didymocarpeae;
                                                                                                                                                                                                                   PubMed=15659624; DOI=10.1105/tpc.104.028936; Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.; "The Role of KNOX Genes in the Evolution of Morphological
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Sukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; lamiids; Lamiales; Gesneriaceae; Cyrtandroideae; Didymocarpeae;
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NCBI_TaxID=167311;
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Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.;
"The Role of KNOX Genes in the Evolution of Morphological Novelty in
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Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.;
"The Role of KNOX Genes in the Evolution of Morphological Novelty
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STM2 protein (Fragment).
Streptocarpus dunnii.
Streptocarpus dunnii.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
Spermatophyta; Magnoliophyta; eudicotyledons; Core eudicots; asterids;
Streptocarpus.
NCBI TaxID=121487;
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PubMed=15659624; DOI=10.1105/tpc.104.028936;

Harrison J., Moller M., Langdale J., Cronk Q., Hudson A.;

"The Role of KNOX Genes in the Evolution of Morphological Novelty in Streptocarpus.";

Plant Cell 17:430-443(2005).

EMBL; AY662103; AAW33757.1; -; Genomic_DNA.

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Majorly a
30 KD pro
Peptide e
OE8 antib
Human OE8
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CC -map) composition. The HMW-map composition was isolated from a D. farinate Created by gel filtration, with each fraction being analysed for the CC presence of proteins that bound to IgE present in mite-allergic dog CC antisera. The HMW-map composition comprises mapA (a 109 kD protein) and CC encoding them, may be used in therapeutic compositions to modify an CC encoding them, may be used in therapeutic compositions to modify an CC encoding them, may be used in therapeutic compositions to modify an CC treated include mammals and birds, especially felines, canines, equines, CC treated include mammals and birds, especially felines, canines, equines, CC treated include mammals and birds, especially felines, canines, equines, CC treated include mammals and birds, especially felines, canines, equines, CC treated include so be used to diagnose allergies via a skin test. The CC proteins and peptides can also be used to raise antibodies, which have a CC variety of potential uses. For example, they can be used as vaccines to CC passitively immunise animals against dust mite hypersensitivity, as CC positive controls in test kits and as tools to recover desired dust mite and allergens from a mixture of proteins
                                                                                                 Matches
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                                                                                                                                                                                              Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequences AAY52510-Y52522 represent proteolytic fragments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Page 69; 154pp; English.
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13-MAY-1998;
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  DPAKGMSPPGFIVGEEGVLS
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98US-0098909P.
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ADO76718
ADO76801
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                                                                                                                   Score 105; DB 3;
Pred. No. 4.7e-09;
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Ado76718 Human :
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RESULT 3
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ID AAU9

AAU96317 standard; peptide; 20

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Dermarophagoides farinae high molecular weight mite allergen protein (HWM composition. The HMW-map composition was isolated from a D. farinae composition was isolated from a D. farinae composition was isolated from a D. farinae composition composition being analysed for the composition of protein being analysed for the composition comprises mapA (a 109 kD protein) and compagnition proteins and peptides, and mucletic acids cencoding them, may be used in therapeutic compositions to modify an analyse created include mammals and birds, especially felines, canines, that may be used in therapeutic compositions to modify an composition to mite allergens. Animals that may be treated include mammals and birds, especially felines, canines, equines, other pets, and work or domestic animals. The proteins or fragments may also be used to diagnose allergies via a skin test. The proteins and peptides can also be used to raise antibodies, which have a variety of potential uses. For example, they can be used as vaccines to positive controls in test kits and as tools to recover desired dust mite allergens from a mixture of proteins
                                                              Matches
                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mite allergen protein; map; high molecular weight; HMW-map; allergy; house dust mite; IgE; immunoglobulin E; allergen; mapA; mapB; hypersensitivity reaction; therapy; treatment; diagnosis; human; feline; canine; veterinary; antibody; vaccine; immunisation.
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13-MAY-1998;
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                DPAKGMSPPGFIVGEEGVLS 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 70; 154pp;
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                                                              Conservative
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                                                                             Score 105; DB 3;
Pred. No. 4.7e-09;
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                                                                Mismatches
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AAU96317;

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                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an isolated mite allergenic protein of C Dermatophagoides, designated Der HMW-map protein, and its related nucleic acid. The Der HMW-map protein is useful for eliciting an immune response against Der HMW-map protein. The protein or a reagent comprising a non-proteinaceous epicope is useful for identifying an animal (e.g., dog, cat) susceptible to or having an allergic response to a mite. An idea is susceptible to or having an allergic response to a mite. The DNA and protein can be used in the detection of anti-Der HMW-map antibodies in animal fluids, and inhibition of immunoglobulin (Ig)E or Der HMW-map protein activity associated with a disease. Antibodies that bind to Der HMW-map are useful for inhibiting binding of proteins to IgE, to prevent immunocomplex formation, thus reducing hypersensitivity responses to mite allergens, and as vaccines against mite allergen hypersensitivity. Sequences AAU96314-AAU96342 represent Der HMW-map polypeptides of the invention
                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 20
                                        Der HMW-map; American house dust mite; antiallergic; mite; IgE; mite allergenic protein; immunoglobulin E; hypersensitivity;
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 Dermatophagoides farinae
                              immunocomplex formation.
                                                                                        Der HMW-map
                                                                                                                        15-JUL-2002
                                                                                                                                                                                    AAU96326 standard; peptide; 20
                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 12; Page 70; 161pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New mite allergenic protein isolated from Dermatophagoides, Der HMW-map protein, useful as a vaccine for treating mite
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-SEP-2000; 2000US-00662293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-SEP-2001; 2001WO-US028730
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Der HMW-map
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-2002
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                                                                                                                                                                                                                                                                                  1 DPAKGMSPPGFIVGEEGVLS 20
                                                                                                                                                                                                                                                                                                                            20;
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                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                              DPAKGMSPPGFIVGEEGVLS 20
                                                                                        polypeptide #13.
                                                                                                                                                                                                                                                                                                                            Conservative
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Pred. No. 4.7e-09;
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RESULT 5
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dermatophagoides, designated Der HMW-map protein, and its related nucleic acid. The Der HMW-map protein is useful for eliciting an immune response against Der HMW-map protein. The protein or a reagent comprising a non-proteinaceous epitope is useful for identifying an animal (e.g., dog, cat) susceptible to or having an allergic response to a mite. A therapeutic composition is useful for desensitising a host animal to an allergic response to a mite. The DNA and protein can be used in the detection of anti-Der HMW-map antibodies in animal fluids, and inhibition of immunoglobulin (Ig)E or Der HMW-map protein activity associated with a disease. Antibodies that bind to Der HMW-map are useful for inhibiting binding of proteins to IgE, to prevent immunocomplex formation, thus reducing hypersensitivity responses to mite allergens, and as vaccines against mite allergen hypersensitivity. Sequences AAU96314-AAU96342 represent Der HMW-map polypeptides of the invention
21-FEB-2003; 2003JP-00045166
21-MAY-2003; 2003JP-00142845
                                              20-FEB-2004; 2004WO-JP002009
                                                                             02-SEP-2004
                                                                                                                                          Murinae
                                                                                                                                                                      antinflammatory; osteopathic.
                                                                                                                                                                                       bone fenestration; bone cartilage transplantation;
                                                                                                                                                                                                                      proliferati
                                                                                                                                                                                                                                     sheathlin; amelin; ameloblastin; enamel matrix protein,
                                                                                                                                                                                                                                                                     Mouse/ rat
                                                                                                                                                                                                                                                                                                    18-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                ADR50710 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-SEP,-2001; 2001WO-US028730
                                                                                                           WO2004074319-A1
                                                                                                                                                                                                    proliferation promoter; differentiation promoter; periodontitis;
fracture; bone defect; osteoporosis; osteoarthritis; bone grafting;
                                                                                                                                                                                                                                                                                                                                    ADR50710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated mite allergenic protein of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 12; Page 71; 161pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New mite allergenic protein isolated from Dermatophagoides, designa
Der HMW-map protein, useful as a vaccine for treating mite allergy.
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                                                                                                                                                                                                                                                                   sheathlin peptide SeqID
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                                                                                                                                                                                                                                                                                                    (first entry)
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RESULT 6
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fracture, bone defect, osteoporosis, osteoarthritis, as well as in bone grafting after surgical removal of tumour, bone fenestration, bone cartilage transplantation and bone supplementation. Accordingly, they exhibit antinflammatory and osteopathic activities. This peptide sequencis a mouse/rat sheathlin peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to novel partial peptides of the sheathlin protein (also known as amelin and ameloblastin), which is an enamel matrix protein Specifically, it refers to physiolocially active peptides that can be employed as promoters of proliferation and differentiation in various cells including osteoblasts, chondroblasts, cementoblasts, bone marrrow-origin mesenchymal stem cells and periodontal membrane cells. The present invention describes peptides that when used as cell growth promoter sequences can be useful for the treatment of periodontitis,
              The
 The present invention relates to modified nitrilase proteins, useful for preparing carboxylic acids, preferably substituted
                                        Disclosure; Page 13; 96pp; English.
                                                                                                                                    Zelinski T,
                                                                                                                                                                                         27-FEB-2003; 2003US-0450470P
                                                                                                                                                                                                                    24-FEB-2004; 2004WO-EP001804
                                                                                                                                                                                                                                               10-SEP-2004
                                                                                                                                                                                                                                                                         WO2004076655-A1
                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                                              Nitrilase; carboxylic acid; enzyme; nitrile
                                                                                                                                                                                                                                                                                                                                                       Nitrilase
                                                                                                                                                                                                                                                                                                                                                                                   02-DEC-2004
                                                                                                                                                                                                                                                                                                                                                                                                              ADR67576;
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADR67576 standard; peptide; 16 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            surgical removal of tumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel partial peptide of sheathlin, useful for treating periodontitis, fracture, osteoporosis, osteoarthritis and useful in bone grafting aft
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Takata T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SEGK ) SEIKAGAKU
                                                                                                                                                                 (BADI )
                                                                  modified nitrilase polypeptides, ds, preferably substituted chiral
                                                                                                            2004-653416/63
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                                                                                                                                                                                                                                                                                                                                                       peptide
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                                                                                                                                     Kesseler M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ID NO 6;
                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                        fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          35.2%;
75.0%;
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                                                                                                                                     Hauer B,
                                                                                                                                                                                                                                                                                                                                                        #165
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Pred. No. 1.8e+02
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                                                                                                                                       Friedrich
                                                                    useful for carboxylic
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                                                                   preparing carboxylic acids.
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chiral
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nitriles into the corresponding carboxylic acids and ammonium ions. The modified nitrilases of the invention have a modification at position 296 and exhibit a modulated acceptance in comparison to wild-type nitrilase. The substitution of tyrosine at position 296 in wild-type Alcaligenes faecalis nitrilase comprising a sequence of 356 amino acids (ADR67405) was realized by site-directed mutagenesis utilizing overlap-extension PCR. The present sequence is a fragment of a nitrilase used in a sequence alignment to illustrate the invention.
                                                                                                                                                                                   Novel CASB88 polypeptide and polynucleotide useful for treating cancer, Crohn's disease, colitis ulcerosa, colorectal cancer, lung cancer and reasons. Wilm's tumor, retinoblastoma and rhabdomyosarcoma.
                                            Sequences of the invention are used for the treatment of diseases e.g. Crohn's disease, Colitis ulcerosa, colorectal cancer, lung cancer and preneoplasic lesions, breast, brain, uterus, muscle, eye and germ cell cancers, Wilm's tumour, retinoblastoma, rabdomyosarcoma, leimyosarcoma and synovial sarcoma. They are useful in gene therapy and as vaccines.
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27-MAR-2001; 2001GB-00007632.
02-AUG-2001; 2001GB-00018926.
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 Sequence 9 AA;
                                                                                                                                                                                                                                                                                Cassart J,
                                                                                                                                                                                                                                                                                                                                                                                                    18-SEP-2001; 2001WO-EP010980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CASB88 protein; Crocolorectal cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human CASB88 antigenic epitope #19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           carboxylic acids. Nitrilases are enzymes which catalyse the hydrolysis of
                                The present sequence is human CASB88
                                                                                                                           The invention relates to novel CASB88 polypeptides and polynucleotides
                                                                                                                                                      Claim 6; Page 77; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                      27-JUN-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rabdomyosarcoma; leimyosarcoma; synovial sarcoma; vaccine; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-DEC-2002
                                                                                                                                                                                                                                                                                                              (SMIK ) SMITHKLINE BEECHAM BIOLOGICALS
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7; Conserv
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                                                                                                                                                                                                                                                                              Coche T,
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Pred. No. 1.9e
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                                antigenic
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1.9e+02;
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                                 epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer; gene therapy;
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Matches Query Match Best Local

Similarity 6; Conserv

Conservative

0,

33.3%;

Score 35; Pred. No. Mismatches

DB 5; 2e+06; 0 Length

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                                                                                                                              Query Match
Best Local &
                                                                                                                                                                    The invention relates to novel CASB88 polypeptides and polynucleotides. Sequences of the invention are used for the treatment of diseases e.g. Crohn's disease, Colitis ulcerosa, colorectal cancer, lung cancer and preneoplasic lesions, breast, brain, uterus, muscle, eye and germ cell cancers, Wilm's tumour, retinoblastoma, rabdomyosarcoma, leimyosarcoma and synovial sarcoma. They are useful in gene therapy and as vaccines. The present sequence is human CASB88 antigenic epitope
                                                                                                                                                                                                                                                    Novel CASB88 polypeptide and polynucleotide useful for treating cancer, Crohn's disease, colitis ulcerosa, colorectal cancer, lung cancer and preneoplasic lesions, Wilm's tumor, retinoblastoma and rhabdomyosarcoma
Cancer-related DGI-5-binder peptide -
                 01-JAN-2004
                                                ADC99464 standard;
                                                                                                                                                      Sequence
                                                                                                                                                                                                                                       Claim 6;
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                                                                                                                                                                                                                                                                                                                                      20-DEC-2000; 2000GB-00031095.
27-MAR-2001; 2001GB-00007632.
02-AUG-2001; 2001GB-00018926.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               rabdomyosarcoma;
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                                                                                                                     Local Similarity
nes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein;
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                                                                                        GMSPPG 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer; Wilm's tumoun rcoma; leimyosarcoma;
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                                                                                                                       Conservative
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                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Crohn's disease; Colitis ulcerosa; preneoplasic lesion; 
ar; Wilm's tumour; retinoblastoma; cancer; gene therapy;
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                                                                                                                                                                                                                                                                                                      ;
                                              peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide;
                                                                                                                              33.3%;
                                                                                                                                                                                                                                                                                                      Gaulis SRJ,
                                                                                                                              Score 35;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               synovial
                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                      Orntoft T,
SEQ ID 302
                                                                                                                      1.6e+02;
thes 0;
                                                                                                                                      DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                              sarcoma;
                                                                                                                                                                                                                                                                                                      Vinals Y De BassolsC;
                                                                                                                                     Length 10;
                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             vaccine; human;
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RESULT 10
AAW65590
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     binder pairs comprising mixing in a reaction vessel phage expressing biological targets and phage expressing target binders, each having distinguishable selection markers and selecting target and target binder pairs based on the selection markers. The molecules of the invention demonstrate cytostatic activity whilst the method may be useful for selecting target and target binder pairs for preparing a composition for treating cancer. Furthermore, the method may be utilised during gene therapy procedures. The current sequence is that of the cancer-related DGI-5-binder peptide of the invention.
16-JUL-1998.
                                                                                                                                       Epstein-Barr virus; diagnostic test; autoimmune infection; antibody; screening; genetic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Selecting target and target binder pairs for preparing a compos
treating cancer by mixing in a reaction vessel phage expressing
biological targets and phage expressing target binders.
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                                                WO9830586-A2
                                                                                              Synthetic.
                                                                                                                                                                                                                    Peptide
                                                                                                                                                                                                                                                                     15-0CT-1998
                                                                                                                                                                                                                                                                                                                     AAW65590;
                                                                                                                                                                                                                                                                                                                                                                   AAW65590 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel method of selecting target and target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 26; SEQ ID NO 302; 172pp; English
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Prendergast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-OCT-2001; 2001US-0345471P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-OCT-2002;
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8; Conservative
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                                                                                                                                                                                                                                                                     (first entry)
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, Goldstein N;
                                                                                                                                                                                                                                                                                                                                                                   peptide;
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Pred. No. 3.4e+02;
1; Mismatches 6;
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                                                                                                                                                                   disease; vaccine;
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13-JAN-1998;

98WO-US000342

97US-00781296

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC virus or viral component to alleviate or prevent the autoimmune disorder. CC Also claimed are: (1) a diagnostic test kit comprising: (a) reagents combined to the virus of experiments of the same to detect levels of antibodies to EBV, indicators of CC which can be used to detect levels of antibodies to EBV, indicators of CC control samples from individuals not at risk of developing an autoimmune disease; and (c) a device for determining the differences in levels of a patient and control samples to distinguish individuals at higher risk of CC developing an autoimmune disease; and (2) a method for screening for genetic markers or risk factors for development of autoimmune disease; and (2) a method for screening for genetic markers or risk factors for development of autoimmune diseases or risk factors for development of autoimmune diseases of different confidence an autoimmune response in at least one of the strains of the same species of an animal vaccinated with EBV or a component to induce an autoimmune response in at least one of the strains of cidentify potential genetic markers or risk factors. The methods can be cused for the prevention, diagnosis, and treatment of autoimmune diseases consisted with EBV as an etiological agent. The autoimmune diseases may be e.g. gystemic lupus erythematosus, Sjogren's syndrome, rheumatoid arthritis, consisted diabetes mellitus, Wegener's granulomatosis, etc. The consisted diseases mellitus, Wegener's granulomatosis, etc. The syndrome, response is shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local (
hypoparathyroidism; primary biliary cirrhosis; ankylosing spondylitis; inflammatory bowel disease; Addison's disease; thyroiditis; filariasis; Graves' disease; ulcerative colitis; dermatcomyositis; myasthenia gravis; Crohn's disease; (CREST syndrome; autoimmune cystitis; glomerulonephritis; polyarteritis nodosa; hepatitis; atopic rhinitis; Goodpasture's syndrome; sarcoidosis; rheumatic fever; anti-phospholipid syndrome; farmer's lung; cushing's syndrome; bird-fancier's lung; alveolitis; erythema nodosum; pyoderma gangrenosum; fibromyalgia; Kawasaki's disease; cardiomyopathy; Sampter's syndrome; asthma; polymyalgia rheumatica; psoriasis; arteritis; erythroblastosis foetalis; cyclitis; IgA nephropathy; Hodgkin's lymphoma;
                                                                                                                                                                                                                                                      polymyositis; systemic lupus erythematosus; SLE; rheumatoid arthritis; Sjogren's syndrome; diabetes mellitus; adrenalitis; multiple sclerosis; demyelinating disease; Hashimoto's thyroiditis; autoimmune infertility;
                                                                                                                                                                                                                                                                                                                                  Vaccine; therapy; autoimmune
                                                                                                                                                                                                                                                                                                                                                                                    Human autoantigen Sm B/B' octapeptide #3 used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                      15-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAE09067 standard; peptide; 8 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Use of Epstein-Barr virus or component(s) - for developing product(s) which can be used for preventing, diagnosing, treating or determining
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprises EBV or a component in a carrier for administration of the
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Pred. No. 2e+0
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                  disorder; Epstein-Barr virus; EBV; anaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
2e+06;
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renal cell carcinoma; eosinophilia; immunosuppressive; ophthalmological; thyromimetic; neuroprotective; cytostatic; nephrotropic; antiallergic; dengue; antiulcer; vasotropic; antipyretic; hepatotropic; human.
                                                                                                                                                                                                                                                                                                                     Novel vaccine for alleviating or preventing autoimmune disorders induced Epstein-Barr virus (EBV) infection e.g. systemic lupus erythematosus, juvenile onset diabetes mellitus, comprises EBV virus or its component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-FEB-2000; 2000US-00500904
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comprising EBV or its component in a carrier. The vaccine is useful for preventing or alleviating autoimmune disorders induced by EBV, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (OKLA-) OKLAHOMA MEDICAL RES FOUND
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cc inflammatory bowel disease, polymyositis, dermatomyositis, multiple condocrine failure, Schmidt's syndrome, autoimmune uveitis, Addison's cc disease, adrenalitis, primary biliary cirrhosis, Graves' disease, ct thyroiditis, Hashimoto's thyroiditis, autoimmune thyroid disease, ct thyroiditis, Hashimoto's thyroiditis, autoimmune thyroid disease, comparisons and haemolytic anaemia, lupoid hepatitis, demyelinating chyporatelyroidism, Dressler's syndrome, myasthenia gravis, autoimmune cystitis, pemphigoid, dermatitis herpetiformis, alopecia areata, comparathyroidism, Dressler's syndrome, myasthenia gravis, autoimmune pemphigus vulgaris, computitis, pemphigoid, dermatitis herpetiformis, alopecia areata, compensione, cystitis, pemphigoid, dermatitis herpetiformis, alopecia areata, compensione, cystitis, pemphigoid, scleroderma, progressive systemic controlactyly and telangiectasia), adult onset diabetes mellitus (Type II conditional pemphigus vulgaris, controlactyly and telangiectasia), adult onset diabetes mellitus (Type II conditional pemphigus), atopic dermatitis, atopic variatity, antylosing spondylitis, collectasia, polyarteritis, atopic dermatitis, atopic variatis, doodpasture's glomyarulonephritis, atopic dermatitis, atopic variatis, foodpasture's glomyarulonephritis, atopic dermatitis, atopic variatis, foodpasture's lung, erythema compensione, cushing's syndrome, autoimmune controlactive hepatitis, bird-fancier's lung, allergic calveditis, toxic necrodermal lysis, alopecia, Alport's syndrome, call arteritis, dengue, cardityle, providerma, cardityle, intersetitial lung disease, contenderma, providerma, encephalomyalia, flavasaus's arteritis, dengue, cardityle, encephalomyositis, intersetitis, and Behcet's contender of lymbomatica, temporal arteritis, shunan's syndrome, largityle, providerme, cardityle, providerme, southerne, logorialia, flavasis, contender of lymbomatic cyclitis, heterochromic cyclitis, heterochromic cyclitis, shundan's syndrome, logorialia, mand bencer's contender of lymbomatical cyclitis, s non-Hodgkin's lymphoma, cardiomyopathy, Henoch-Schonlein purpura, post vaccination syndromes, renal cell carcinoma, Eaton-Lambert syndrome or systemic lupus erythematosus (SLE), Sjogren's syndrome, juvenile onset diabetes mellitus, rheumatoid arthritis, Wegener's granulomatosis, The present invention relates to a vaccine for alleviating or preventing autoimmune disorders induced by infection with Epstein-Barr virus (EBV), polychondritis. used in the The present sequence is human autoantigen Sm The invention invention

Query Match Best Local Matches l Similarity 6; Conserv Conservative 32.4%; Score 34; DB 4; Pred. No. 2e+06; DB 4; 2 Length 8 0 Gaps

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PAKGMSPP

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RESULT 12
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AX AAB97
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                                                                                    This invention relates to a B cell clonal toxin. The toxin is made from CC two moleties, the first causes the toxin to be internalised by a B cell, CC and the second is a biologically acceptable toxin. The invention includes CC amethod for inactivating/killing an antigen specific B cell. A target B CC cell is contacted with an effective immunosuppression in conditions CC method is useful for selective immunosuppression in conditions CC characterised by the presence of an unwanted or deleterious immune CC disease conditions. Use of the B cell clonal toxin can result in CC immunosuppressive; antiinflammatory; antiallargic; virucide; antidiabetic cophthalmological; and nephrotropic; cardiant; antiulcer; cophthalmological; and nephrotropic activity. The toxin is particularly CC useful for treating a host suffering from an antigen specific antibody condition, where the antigen specific antibody is CC useful for treating a host suffering from an antigen specific antibody condition useful for treating allergies, viral disease conditions, and condomnune disorders. Also treated are skin disease; autoimmune cCC undoundamentologic disorders; gastrointestinal disease; neurologic and inferttlity disorders. The present as skin disease; pulmonary diseases; and inferttlity disorders. The present activity all seases; pulmonary diseases; and inferttlity disorders. The present sequence represents all RMP with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B cell clonal toxin useful for treating autoimmune disorders such Grave's disease, myocardial infarction, Crohn's disease, multiple sclerosis, comprises a group that causes toxin to be internalized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunosuppressive; antiinflammatory; antiallergic; antidiabetic; thyromimetic; antithyroid, vasotropic; cardiant; antiulcer; neuroprotective; antirheumatic; antiarthritic; dermatological; ophthalmological; nephrotropic; antiarthritic; dermatological; ophthalmological; nephrotropic; allergy; autoimmune disorder; skin disease; autoimmune endocrinopathy; vasculitic syndrome; cardiovascular disease; immunohaematologic disorder; neurologic disease; gastrointestinal disease; collagen vascular disease; renal disease; pulmonary disease; infertility disorder; UI-U6 RNP; systemic lupus erythetmatosus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 33; 46pp; English
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infertility disorders. The present sequence represents a UI RNP with nd C proteins epitope. An antibody response to this antigen is licated in systemic lupus erythematosus, a disorder which may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Н
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comprising an Epstein-Barr virus or a component in a carrier for administration to alleviate or prevent the autoimmune discorders. The methods and compositions of the present invention are useful for compositions of the present invention are useful for control of the preventing autoimmune discorders, such as diabetes mellitus, rheumatoid arthritis, juvenile rheumatoid carrhritis, osteoarthritis, psoriatic arthritis, multiple sclerosis, control of the providitis, active carrier to the screen control of the sclerosis, control of the screen control of the sclerosis, control of the screen co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 13
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16-MAY-1996;
13-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a vaccine for alleviating or preventing autoimmune disorders induced by infection with Epstein-Barr virus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New vaccine preventing or alleviating autoimmune disorders induced by the 
Epstein-Barr virus, such as diabetes mellitus, rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Harley JB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-OCT-2001; 2001US-00012756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2002164355-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Epstein-Barr virus nuclear antigen peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-OCT-2003
10-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treated using the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HARL/) HARLEY J B. (JAME/) JAMES J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EBV; viral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU07632 standard; peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sclerosis, systemic lupus erythematosus, atopic dermatitis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PAKGMSPP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .; spstein-Barr virus nuclear antigen; vaccine; disorder; Epstein-Barr virus infection; diabetes mellitus; multiple sclerosis; dermatitis; psoriasis; asthma; anaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          James JA
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fig 3; 41pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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(first entry)
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96US-0019053P.
97US-00781296.
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Pred. No. 2e+06;
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                                                              organs or organ systems, vascular stenosis, occlusion to peripheral vessels (such as limb), hypertension, and on the treatment of a conditic or disorder that can be ameliorated with the selective bradykinin receptor agonist. Also useful for treatment of a condition for which dilation of vessels is required e.g. treatment of tumours and cancers, and for gene therapy. Maximakinin is a selective bradykinin receptor agonist and exhibits both tissue and receptor specific targeting. This tissue selectivity enables the use of the peptides in the treatment of diseases of the cardiovascular system while minimizing the side effects associated with the use of less selective agents. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a bradykinin agonist, useful in the selective stimulation of arterial smooth muscle bradykinin receptors or treatment of cardiovascular disease comprises maximakinin, its analogue or fragment or their encoding nucleic acids. Maximakinin derivatives are used in the preparation of a medicament for selectively stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bradykinin agonists comprising maximakinin and its fragments, useful selectively stimulating arterial smooth muscle bradykinin receptors of treating cardiovascular disease e.g. ischemic heart disease.
                                                                                                                                                                                                                                                                                                                                                                                    arterial smooth muscle bradykinin receptors; for treating cardiovascular disease (claimed) e.g. ischemic heart disease, ischemic disease of other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 7;
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06-JUN-2003;
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                                    represents a maximakinin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UUTE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 3; 67pp; English.
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2e+06;
                                    bradykinin agonist
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Matches Best Local Query Match

6. Similarity

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                        organs or organ systems, vascular stenosis, occlusion to peripheral vessels (such as limb), hypertension; and on the treatment of a condition or disorder that can be ameliorated with the selective bradykinin receptor agonist. Also useful for treatment of a condition for which dilation of vessels is required e.g. treatment of a condition for which agonist and exhibits both tissue and receptor specific targeting. This tissue selectivity enables the use of the peptides in the treatment of diseases of the cardiovascular system while minimizing the side effects associated with the use of less selective agents. The present sequence represents a maximakinin derivative as a bradykinin agonist
                                                                                                                                                                                                  The present invention relates to a bradykinin agonist, useful in the selective stimulation of arterial smooth muscle bradykinin receptors or treatment of cardiovascular disease comprises maximakinin, its analogue or fragment or their encoding nucleic acids. Maximakinin derivatives are used in the preparation of a medicament for selectively stimulating arterial smooth muscle bradykinin receptors; for treating cardiovascular disease (claimed) e.g. ischemic heart disease, ischemic disease of other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 11
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Sequence 12
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06-JUN-2003; 2003GB-00012992.
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vascular stenosis; hypertension; tumour; cancer; Maximakinin.
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                                                                                                                                                                                                                                                                                                                                   67pp; English.
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Pred. No. 2.
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RESULT 17
ABP48137
ID ABP48
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                                                                                                                                               Query Match
Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                              organs or organ systems, vascular stenosis, occlusion to peripheral vessels (such as limb), hypertension; and on the treatment of a condition or disorder that can be ameliorated with the selective bradykinin receptor agonist. Also useful for treatment of a condition for which dilation of vessels is required e.g. treatment of tumours and cancers, and for gene therapy. Maximakinin is a selective bradykinin receptor agonist and exhibits both tissue and receptor specific targeting. This diseases of the cardiovascular system while minimizing the side effects associated with the use of less selective agents. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a bradykinin agonist, useful in the selective stimulation of arterial smooth muscle bradykinin receptors or treatment of cardiovascular disease comprises maximakinin, its analogue or fragment or their encoding nucleic acids. Maximakinin derivatives are used in the preparation of a medicament for selectively stimulating arterial smooth muscle bradykinin receptors; for treating cardiovascular disease (claimed) e.g. ischemic heart disease, ischemic disease of other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bradykinin agonists comprising maximakinin and its fragments, useful for selectively stimulating arterial smooth muscle bradykinin receptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-FEB-2003; 2003GB-00002623
06-JUN-2003; 2003GB-00012992
ABP48137 standard; peptide; 18
                                                                                                                                                                                                                Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 7; SEQ ID NO 6; 67pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               bradykinin; Cardiovascular; Vasotropic; Hypotensive; Cytostatic; vascular stenosis; hypertension; tumour; cancer; Maximakinin.
                                                                                                                                                                                                                                                  represents a maximakinin derivative as a bradykinin agonist
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Pred. No. 3.1e+02;
0; Mismatches 2;
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RESULT 18
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Best Local (
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06-FEB-2003; 2003GB-00002623
06-JUN-2003; 2003GB-00012992
                                                                                 WO2004069857-A2
                                     06-FEB-2004; 2004WO-GB000470
                                                                                                                                  bradykinin; Cardiovascular; Vasotropic; Hypotensive; Cytostatic;
vascular stenosis; hypertension; tumour; cancer; Maximakinin.
                                                                                                                                                                                                  04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 AA;
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                                                             19-AUG-2004.
                                                                                                             Unidentified
                                                                                                                                                                       Maximakinin 1-18 as a bradykinin agonist.
                                                                                                                                                                                                                                                  ADR43748 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Fig 15A; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hormones and are useful to treat conditions such as dwarfism a acromegaly, and to promote growth and milk production in cows.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New amino acid sequences fully defined in the specification mimic hormones and are useful to treat conditions such as dwarfism and
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6; Conserve
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                                                                                                                                                                                                                                                 peptide;
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Pred. No. 4.4e+02;
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RESULT 19
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a bradykinin agonist, useful in the selective stimulation of arterial smooth muscle bradykinin receptors or treatment of cardiovascular disease comprises maximakinin, its analogue or fragment or their encoding nucleic acids. Maximakinin derivatives are used in the preparation of a medicament for selectively stimulating arterial smooth muscle bradykinin receptors; for treating cardiovascular disease (claimed) e.g. ischemic heart disease, ischemic disease of other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bradykinin agonists comprising maximakinin and its fragments, useful foselectively stimulating arterial smooth muscle bradykinin receptors and for treating cardiovascular disease e.g. ischemic heart disease.
             WPI; 2004-365151/34
                                                                                                                                                                                                     WO2004035614-A1
                                                                                                                                                                                                                                 Unidentified
                                                                                                                                                                                                                                                            allosteric modulator; g-alpha
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                                                                                                                                                                                                                                                                                                                                                                                           ADN17072 standard; peptide;
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                                          Hyde-Deruyscher R,
                                                                                                               18-OCT-2002; 2002US-0419143P
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                                           Duffin
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75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 34; DB 8;
Pred. No. 4.4e+02;
                                          Hamilton PT, Fredericks
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RESULT 20
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Best Local
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                                                                                                                                               Bradykinin agonists comprising maximakinin and its fragments, useful selectively stimulating arterial smooth muscle bradykinin receptors after treating cardiovascular disease e.g. ischemic heart disease.
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                                                                                           Disclosure;
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06-JUN-2003; 2003GB-00012992
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8; Conserv
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                                                                                                                                                                                                                                                                                                                              Hirst D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cardiovascular; Vasotropic; Hypotensive; Cytostatic; enosis; hypertension; tumour; cancer; Maximakinin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA,
                                                                                           SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                              ö
                                                                                                                                                                                                                                                                                                                                 Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           32.4%;
                                                                                        1; 67pp; English
                                                                                                                                                                                                                                                                                                                     T,
                                                                                                                                                                                                                                                                                                                                 O'rourke
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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The present invention

relates to a bradykinin agonist, useful in the of arterial smooth muscle bradykinin receptors

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selective stimulation

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RESULT 21
AAW18175
ID AAW18
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Best Local
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20-OCT-1995;
31-OCT-1995;
06-DEC-1995;
A vaccinating agent for promoting an immune response in a mammal against Mycobacterium pathogens comprises at least one immunodominant epitope of at least one majorily abundant extracellular protein, i.e. the M.tuberculosis 110, 80, 71, 58, 45, 32A, 32B, 30, 24, 23.5, 23, 16, 14 or 12 kD proteins, or their analogues, homologues and subunits. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 arterial smooth muscle bradykinin receptors; for treating cardiovascular disease (claimed) e.g. ischemic heart disease, ischemic disease of other organs or organ systems, vascular stenosis, occlusion to peripheral vessels (such as limb), hypertension; and on the treatment of a condition or disorder that can be ameliorated with the selective bradykinin receptor agonist. Also useful for treatment of a condition for which dilation of vessels is required e.g. treatment of tumours and cancers, and for gene therapy. Maximakinin is a selective bradykinin receptor agonist and exhibits both tissue and receptor specific targeting. This tissue selectivity enables the use of the peptides in the treatment of diseases of the cardiovascular system while minimizing the side effects associated with the use of less selective agents. The present sequence represents maximakinin 1-19 as a bradykinin agonist
                                                                                                                                                                                                                                           Vaccines derived from M.tuberculosis major abundant extracellular proteins - are easy to prepare and less toxic than conventional killed attenuated vaccines, useful for protecting against or treating
                                                                                                                                                                  Claim 49; Page 162; 193pp;
                                                                                                                                                                                                                             Mycobacterial infections.
                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-020936/02.
                                                                                                                                                                                                                                                                                                                                                                                                                 Horwitz MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAY-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunodominant epitope from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (REGC )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VIND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protozoan;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tuberculosis
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95US-00545926.
95US-00551149.
95US-00568357.
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                                                                                                                                                                  English.
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Pred. No.
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2;
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Best Local
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         Sequences shown in AAW75587 to AAW75641 represent synthetic peptides derived from the native 30kD major secretory protein of M. tuberculosis. These peptides are used for splenic lymphocyte profileration assays to identify the immunodominant T-cell epitope of the 30kD protein. The invention provides an agent for vaccinating mammals against Mycobacterium. The agent comprises at least one of the major abundant Mycobacterium agent comprises at least one of the major abundant extracellular 110, 80, 71, 58, 45, 32A, 32B, 30, 24, 23.5, 23, 16, 14 or 12 kD proteins of M. tuberculosis, or at least 1 of their immunodominant epitopes and interleukin-12 (IL-12) or MF59 as adjuvants. The agent
                                                                                                                                                                                                                             Vaccines against Mycobacterium containing major extracellular proteins used to, e.g. induce protective and therapeutic immune responses, and f detecting an immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vectors, containing the DNA for the extracellular proteins, are used to transform cells for production of recombinant DNA molecules. More generally the DNA from other pathogens can be used in vaccines, e.g. against other bacteria, viruses, fungi and protozoa. Since different combinations of DNA can be used, a wide range of effective compositions can be produced. They generate a response against the antigens most often found on infected cells during the infection, regardless of the strength or specificity of the immune response. The vaccines are easy to produce and less toxic than known killed or attenuated vaccines, so can be given to immunocompromised subjects, e.g. those with HIV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mycobacterium tuberculosis; vaccination; extracellular product; immunodominant epitope; interleukin-12; MF59; immune response; opsonising humoral response; intracellular pathogen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      M. tuberculosis 30
                                                                                                                                                                                           Example 28; Page
                                                                                                                                                                                                                                                                                                                                        Horwitz MA,
                                                                                                                                                                                                                                                                                                                                                                                                               21-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protect against (or to treat existing) infections by Mycobacterium (especially M. tuberculosis) while the epitopes can also be used to detect presence of an immune response to a Mycobacterium pathogen.
                                                                                                                                                                                                                                                                                                                                                                              (REGC ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mycobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           extracellular product 30 KD protein.
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                                                                                                                                                                                                                                                                                                                                                                              CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tuberculosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US000942
                                                                                                                                                                                           96; 236pp; English.
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                                                                                                                                                                                                                                                                                                                                          Lee
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Pred. No. 5.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        derived
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31 (residues 151-165).
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RESULT 23
ABU56381
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                          The invention describes a vaccine for promoting an immune response, in a mammalian host, against infectious pathogens of the genus Mycobacterium, comprising at least 1 immunodominant epitope of at least one majority abundant extracellular product comprising Mycobacterium tuberculosis 110, 80, 71, 58, 45, 32A, 32B, 30, 24, 23.5, 23, 16, 14, 12 KD protein or their analogues, homologues or subunits. The proteins and polypeptides of the invention are useful in gene therapy and treatment of diseases caused by Mycobacterium such as tuberculosis. This sequence represents a
                                                                                                                                                                                                                                                                                                                                                  06-DEC-1995;
23-MAY-1996;
21-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                             23-NOV-1993;
12-AUG-1994;
23-MAY-1995;
31-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mycobacterium, specifically M. tuberculosis. The immunodominant epitopes can also be used (typically in a cutaneous hypersensitivity test) to detect an immune response to vaccination. Preparation of the agent does not require selection of the most immunogenic products, so large scale production and purification are easy, resulting in a consistent, standardised formulation, having lower toxicity than killed or attenuated vaccines. The agents provide a rapid and effective response (including a strong cell-mediated component) and are safe even in immunocompromised subjects. They prevent development of an opsonising humoral response that might spread intracellular pathogens
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Sequence 15
                                                                                                                                                                                                                              New vaccine,
                                                                                                                                                                                                                                                                                          Horwitz
                                                                                                                                                                                                                                                                                                                      (REGC ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-SEP-2001; 2001US-00953510.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2002131975-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antibacterial; tuberculosis; vaccine; gene therapy; 30kDa protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mycobacterium tuberculosis 30kDa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-MAR-2003
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                             fragment of the Mycobacterium tuberculosis 30kDa protein
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                                                                                                                                                                                  Page 66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DPSQGMGP 11
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                                                                                                                                                                                                             useful for promoting an immune response againthe genus Mycobacterium in a mammalian host.
                                                                                                                                                                                                                                                                                        Harth G;
                                                                                                                                                                                                                                                                                                                      CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                 94US-00289667.
95US-00447398.
95US-00551149.
95US-00568357.
96US-00552842.
98US-00157689.
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                                                                                                                                                                                 82pp; English
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12-AUG-1994;
23-MAY-1995;
31-OCT-1995;
06-DEC-1995;
23-MAY-1996;
                                                                                                                                                                                                                                                        The invention relates to a novel nucleic acid encoding an abundant extracellular protein of Mycobacterium tuberculosis useful as vaccines for generating protective or therapeutic immune response against viral, bacterial, fungal and protozoal infections. They are also used as immunotherapeutic agents. The present sequence is M. tuberculosis strain Erdman 30 KD major secretory protein immunodominant T-cell epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A nucleic acid encoding an abundant extracellular protein of Mycobacterium tuberculosis useful as vaccines for generating or therapeutic immune response against viral, bacterial, fund
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vaccine; a virucide;
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                                                                                                                                                                                        Sequence
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DPSQGMGP 11
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Pred. No. 5.1e+02;
2; Mismatches 1
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RESULT 25 ADF45168 ID ADF45

ADF45168 standard; peptide; 15

ADF45168;

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Search completed: January 26, 2006, 07:58:12 Job time : 70.931 secs
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                                                                                                            Query Match 31.4
Best Local Similarity 62.9
Matches 5; Conservative
                                                                                                                                                                                                          The invention describes a new vaccinating agent for promoting an immune response in a mammalian host against Mycobacterium comprising at least one immunodominant epitope of at least one majorly abundant extracellular product comprising M. tuberculosis 110, 80, 71, 58, 45, 32A, 32B, 30, 24, 23.5, 23, 16, 14 or 12 KD protein, or its analogue, homologue or subunit. The bacterium is M. tuberculosis. The vaccinating agent is useful for promoting an immune response in a mammalian host against Mycobacterium. This is the amino acid sequence of a Mycobacterium tuberculosis 30kD extracellular protein fragment that can be used to illicit and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-AUG-1994
23-MAY-1995
31-OCT-1995
31-OCT-1996
23-MAY-1996
06-DEC-1996
21-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                          Vaccinating agent useful for promoting an immune response in a mammalian host against Mycobacterium, comprises an immunodominant epitope of a majorly abundant extracellular product comprising an M. tuberculosis 30 KD protein.
                                                                                                                                                                        Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                           Claim 3; SEQ ID NO 67; 83pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Horwitz MA;
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Mycobacterium; 30kD; extracellular protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HORW/) HORWITZ M
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95US-00551149.
96US-005632842.
96US-00568357.
98US-00157689.
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Pred. No. 5.1e+02;
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

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    572060 seqs,
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US-08-99-08-88-28
US-09-657-887-397
US-10-198-053-397
US-10-198-053-397
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US-09-827-271-397
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                                                                                                                    SEQ ID NO 13
LENGTH: 20
TYPE: PRT
ORGANISM: Dermatophagoides farinae
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No. 6455686
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LENGTH: 20
TYPE: PRT
ORGANISM: Dermatophagoides farinae
S-09-292-225-4
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    Query Match
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APPLICANT: McCall, Catherine A.
APPLICANT: Hunter, Shirley Wu
APPLICANT: Weber, Eric R.
TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, PROTEINS
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: AL-2-C3
CURRENT APPLICATION NUMBER: US/09/292,225
CURRENT FILING DATE: 1999-04-15
EARLIER APPLICATION NUMBER: 60/098,909
EARLIER FILING DATE: 1998-09-02
EARLIER FILING DATE: 1998-09-02
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                                                                                                                                                                                                                                                                                                                                                         EARLIER APPLICATION NUMBER: 60/085,295
EARLIER FILING DATE: 1998-05-13
EARLIER FPLICATION NUMBER: 60/098,565
EARLIER FILING DATE: 1998-04-17
EARLIER APPLICATION NUMBER: 09/062,013
EARLIER FILING DATE: 1998-04-17
NUMBER OF SEQ ID NOS: 49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 DPAKGMSPPGFIVGEEGVLS 20
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    100.0%; Score 105;
    DB 2;
Length 20;
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                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,955
FILING DATE: June 7, 1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6641813
GENERAL INFORMATION:
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Matches 6; Conservative
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TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094

CURRENT APPLICATION NUMBER: US/09/270,767

CURRENT FILING DATE: 1999-03-17

NUMBER OF SEQ ID NOS: 62517

SOFTWARE: PatentIn Ver. 2.0
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                                                  FILING DATE: January 31, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/472,94
FILING DATE: January 31, 1990
                                                                                                                                              CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
               ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 2800 O
CITY: Atlanta
STATE: GA
                                                                                                                                                                                 APPLICATION NUMBER: 07/867, FILING DATE: April 13, 1992
                                                                                                                             APPLICATION NUMBER: 07/648,205
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11
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                                                                      07/472,947
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Pred. No.
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); Mismatches 0;
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Query Match
Best Local Similarity
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; ORGANISM: homo sapien
US-07-867-819D-116
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                    Sequence 100, Application US/08475955 Patent No. 6641813 GENERAL INFORMATION:
APPLICANT: Harley, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Pat
SEQ ID NO 116
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Best Local Similarity 75.0%;
Matches 6; Conservative
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TITLE OF INVENTION: Methods and Reagents for Diagnosis of Autoantibodies
FILE REFERENCE: OMRF 114 CIP (2)
CURRENT APPLICATION NUMBER: US/07/867,819D
CURRENT FILING DATE: 1992-04-13
PRIOR APPLICATION NUMBER: 07/472,947
PRIOR APPLICATION NUMBER: 07/472,947
PRIOR FILING DATE: 1990-01-31
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1991-01-31
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                                                                                                                                                           APPLICANT: Harley, John
TITLE OF INVENTION: METHODS AN
TITLE OF INVENTION: AUTOANTIBO
NUMBER OF SEQUENCES: 218
CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                               STATE:
                                                                                                                                          STREET:
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TOPOLOGY: lir
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                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
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                                                                                                                            Atlanta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PatentIn version 3.1
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                                                                                                                                          2800 One Atlantic Center, 1250 West Peachtree Street
                                                                                          USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                 METHODS AND REAGENTS FOR DIAGNOSIS OF AUTOANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32.4%;
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Pred. No. 4.6e+05;
0; Mismatches 2
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Pred. No. 4.6e+05;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 8;
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US-07-867-819D-100
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APPLICANT: Harley, John
TITLE OF INVENTION: Methods and Reagents for Diagnosis of Autoantibodies
FILE REFERENCE: OMRF 114 CIP (2)
CURRENT APPLICATION NUMBER: US/07/867,819D
CURRENT FILING DATE: 1992-04-13
PRIOR APPLICATION NUMBER: 07/472,947
PRIOR FILING DATE: 1990-01-31
PRIOR FILING DATE: 1990-01-31
PRIOR APPLICATION NUMBER: 07/648,205
PRIOR APPLICATION NUMBER: 07/648,205
PRIOR FILING DATE: 1991-01-31
NUMBER OF SEQ ID NOS: 161
NUMBER OF SEQ ID NOS: 161
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 100, Application US/07867819D Patent No. 6897287 GENERAL INFORMATION:
                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1 SEQ ID NO 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 32.4%;
Best Local Similarity 75.0%;
Best Local Similarity 75. Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (404)-873-879
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
NAME: Pabet, Patrea L,
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMFF11.
TELECOMMUNICATION INFORMATION:
TELECHONE: (4041-873-8794
                                                                                       NAME/KEY: MISC FEATURE
LOCATION: (1)...(8)
OTHER INFORMATION: Binding site
                                                                                                                                                     TYPE: PRT
ORGANISM: homo sapien
FEATURE:
                                                                                                                                                                                                             ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
. APPLICATION NUMBER: US/08/475,955
FILING DATE: June 7, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Binding-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6; Conservative
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April 13, 1992
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               32.4%;
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Score 34; DB 2;
Pred. No. 4.6e+05;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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Pred. No. 4.6e+0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .6e+05;
                                   Length 9;
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RESULT 9
US-09-157-689-67
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US-08-990-888-28
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SEQ ID NO 28
LENGTH: 18
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 2598-4002
CURRENT APPLICATION NUMBER: US/08/990,888B
CURRENT FILING DATE: 1997-12-15
NUMBER OF SEQ ID NOS: 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Blume, Arthur J.
APPLICANT: Brissette, Renee
APPLICANT: Carcamo, Juan
APPLICANT: Mandecki, Wlodeck S.
APPLICANT: Tang, Pauline M.
TITLE OF INVENTION: Assays For Compounds Which Bind Growth Hormone Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
           APPLICATION NUMBER: US 08/
FILING DATE: 06-DEC-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 31-OCT-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: H5 peptide secondary library sequence
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                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Abundant Ext
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Horwitz, Marcus A. APPLICANT: Harth, Gunter
                                                                                                                                                                                                                                                                                                                                                              CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
STREET: 20
                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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5. 6387879
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2029 Century Park East, Suite 3800
                                                                                                                                                                                                                                                                                                                                               U.S.A.
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23-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Abundant Extracellular Products and Methods for Their Production and Us
                                                                                                                         US 08/568,357
             US 08/447,398
                                                                  US 08/551,149
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; FRAGMENT TYPE: int
; ORIGINAL SOURCE:
; ORGANISM: Mycoba
; STRAIN: Erdman
US-09-157-689-67
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PATENT NO. 6818223
GENERAL INFORMATION:
APPLICANT: HOTWITZ, Marcus A.
TITLE OF INVENTION: Abundant Extracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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HYPOTHETICAL: N
ANTI-SENSE: NO
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FILING DATE: 23-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 01
FILING DATE: 12-AUG-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
08/289,667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (310) 788-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: MacLean, Kurt A. REGISTRATION NUMBER: 31,118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH:
                                                                                                                                                                              Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/953,510
FILING DATE: 14-Sep-2001
                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Kurt A. MacLean
STREET: 2029 Century Park East, Suite 3800
                            APPLICATION NUMBER: US 08/156,358 FILING DATE: 23-NOV-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                       ZIP: 90067
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                APPLICATION NUMBER: US 08/447,398
FILING DATE: 23-WAY-1995
APPLICATION NUMBER: US 08/289,667
FILING DATE: 12-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
                                                                                                                                                                       CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
REGISTRATION NUMBER: 31,118
                 NAME: MacLean,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15 amino acids
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NO
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CURRENT APPLICATION NUMBER: US/09/667,857
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 455
SOFTWARE: FREESEQ for Windows Version 3.0
SEQ ID NO 397
LENGTH: 20
TYPE: PRT
ORGANISM: Homo sapiens
US-09-667-857-397
                                                                                                                                                                                                                             Query Match
Best Local Similarity
"""hes 5; Conserva
                                                                                                     RESULT 12
US-10-198-053-397
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US-09-667-857-397
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                                                  Sequence 397, Applica Patent No. 6858710 GENERAL INFORMATION:
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APPLICANT: Bangur, Chaitanya S. APPLICANT: Retter, Marc W. APPLICANT: Fanger, Gary R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Retter, Marc W.
APPLICANT: Renger, Gary Richard
APPLICANT: Reed, Steven G.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Mitcham, Jennifer APPLICANT: King, Gordon E. APPLICANT: Algate, Paul A. APPLICANT: Fling, Steven P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 210121.462C5
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INFORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ANTI-SENSE: NO FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: peptide HYPOTHETICAL: NO
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                                                                                                                                                                        13 VGEEGVLS 20
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3 IGEDGILS 10
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TYPE: amino acid
STRANDEDNESS: <Unknown>
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                                                                                       Application US/10198053
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                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jennifer L.
                                                                                                                                                                                                                                                              62.5%;
                                                                                                                                                                                                                                                              Score 33; DB Pred. No. 99;
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CURRENT APPLICATION UNMBER: US/10/198,053
CURRENT FILING DATE: 2002-07-17
NUMBER OF SEQ ID NOS: 624
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 603
LENGTH: 20
TYPE: PRT
ORGANISM: Homo sapiens
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-827-271-397
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US-09-827-271-397
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US-10-198-053-603
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-198-053-397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-198-053-603
                                                                                                                             GENERAL INFORMATION:
APPLICANT: Retter, Marc W.
APPLICANT: Fanger, Gary R.
APPLICANT: FANGER, GARY R.
APPLICANT: FANGER, GARY R.
APPLICANT: FANGER, GARY R.
APPLICANTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121,462C6
CURRENT APPLICATION NUMBER: US/09/827,271
CURRENT FILING DATE: 2001-04-04
                                                   SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 397
LENGTH: 20
                                                                                                                                                                                                                                                                                                    Sequence 397, Appl
Patent No. 6962980
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CURRENT APPLICATION NUMBER: US/10/198,053
CURRENT FILING DATE: 2002-07-17
NUMBER OF SEQ ID NOS: 624
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 31.4%; Score 33; DB Best Local Similarity 62.5%; Pred. No. 99; Matches 5; Conservative 3; Mismatches
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Patent No. 6858710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Retter, Marc W.
APPLICANT: Fanger, Gary R.
APPLICANT: Hill, Paul
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C9
                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bangur, Chaitanya S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hill, Paul
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Similarity 62.5%;
5; Conservative
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Pred. No. 99;
3; Mismatches
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RESULT 16
US-08-478-572-185
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US-08-482-142-185
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; Sequence 185, Application US/08478572
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FRAGMENT TYPE:
US-08-482-142-185
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Best Local Similarity
Matches 5; Conserv
                                                                                                                                                Matches
                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/445,307
FILING DATE: 07 June 1995
ATTORNEY/AGENT INFORMATION:
NAME: CRAIG, ANNE I.
REGISTRATION NUMBER: 32,976
REGISTRATION NUMBER: 32,976
                                                                                                                                                                                                                                                                                           TELEFAX: (617) 466-6040
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,142
FILING DATE: 07-UN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Chen, Ala..
APPLICANT: Evans, Sean
APPLICANT: Evans, Sean
APPLICANT: Shaked, Ze'ev
APPLICANT: Shaked, Ze'ev
TITLE OF INVENTION: T CELL EPITOPES OF THE MAJOR ALLERGENS
TITLE OF INVENTION: FROM DERMATOPHAGOIDES (HOUSE DUST MITE)
NUMBER OF SEQUENCES: 207
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
NORRESSEE: IMMULOGIC PHARMACEUTICAL CORPORATION
NORRESSEE: IMMULOGIC PHARMACEUTICAL CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 466-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                              TYPE: amino acid TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
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ilarity 62.5%;
Conservative
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ilarity 55.6%;
Conservative
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Kuo, Mei-chang
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Pred. No. 99;
3; Mismatches
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Pred. No. 1.6e+02;
4; Mismatches 0;
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RESULT 17
US-08-484-296-185
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Patent No. 626849
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                Query Match 29.5%;
Best Local Similarity 55.6%;
Matches 5; Conservative
                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Garman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617) 466-6040 INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/478,572
FILING DATE: 07-June-1995
CLASSIFICATION:
                                                                                      APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: CRAIG, ANNE I
REGISTRATION NUMBER: 32,976
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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APPLICATION NUMBER: 08/445,307
FILING DATE:
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ADDRESSEE: IMMULOGIC PHARMACEUTICAL CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Evans, Sean
APPLICANT: Shaked, Ze-ev
TITLE OF INVENTION: T CELL EPITOPES OF THE MAJOR ALLERGENS
TITLE OF INVENTION: FROM DERMATOPHAGOIDES (HOUSE DUST MITE)
                APPLICANT: Evans, Sean
APPLICANT: Shaked, Ze'ev
TITLE OF INVENTION: FROM DERMATOPHAGOIDES (HOUSE DUST MITE)
                                                                                                                                             APPLICANT:
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NUMBER OF SEQUENCES:
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CITY: WALTHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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Franzen, Henry
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Kuo, Mei-chang
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Greenstein, Julia
                                                                                        Chen, Xian
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Chen, Xian
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                                                                                                                                                                                   Garman, Richard
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CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMULOGIC PHARMACEUTICAL CORPORATION
STREET: 610 LINCOLN STREET

STREET: 610 LII CITY: WALTHAM STATE: MA

USA

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                Query Match
Best Local Similarity
""" has 5; Conserve
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                                                                                                                   ; FEATURE:
; OTHER INFORMATION: H5 peptide secondary library sequence US-08-990-888-36
                                                                                                                                                                                                                 FILE REFERENCE: 2598-4002

CURRENT APPLICATION NUMBER: US/08/990,888B

CURRENT FILING DATE: 1997-12-15

NUMBER OF SEQ ID NOS: 81

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 36

LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 36, Application US/08990888B Patent No. 6387879 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 29.5%;
Best Local Similarity 55.6%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                     APPLICANT: Blume, Arthur J.
APPLICANT: Brissette, Renee
APPLICANT: Carcamo, Juan
APPLICANT: Mandecki, Wlodeck S.
APPLICANT: Tang, Pauline M.
TITLE OF INVENTION: Assays For Compounds Which Bind Growth Hormone Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/445,307
FILING DATE: 07 June 1995
ATTORNEY/AGENT INFORMATION:
NAME: CRAIG, ANNE I.
REGISTRATION NUMBER: 32,976
REFERENCE/DOCKET NUMBER: 017.6US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 466-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII TEXT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,296
FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (617) 466-6040 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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  5 GMSPPGFIVG 14
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amino acid
OGY: linear
                                     Conservative
                                                       29.5%;
50.0%;
                                     4; Mismatches
                                                     Score 31; DB 2;
Pred. No. 1.9e+02;
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                                     1; Indels
                                                                        Length 18;
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US-09-635-501-29
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US-09-570-022-13
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                                                                Query Match
Best Local S
Matches 6
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APPLICANT: GORDON, R.
APPLICANT: MOORAD,
APPLICANT: DCCTOR,
APPLICANT: GARCIA,
APPLICANT: GARCIA,
                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/40
PRIOR FILING DATE: 1999-09-29
PRIOR APPLICATION NUMBER: 09/16
PRIOR FILING DATE: 1998-09-30
PRIOR APPLICATION NUMBER: 08/98
PRIOR FILING DATE: 1997-12-11
NUMBER OF SEQ ID NOS: 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: ACTON, SUBBAN L. et al.
APPLICANT: ACTON, SUBBAN L. et al.
TITLE OF INVENTION: ANGIOTENSIN CONVERTING ENZYME HOMOLOG AND THERAPEUTIC
TITLE OF INVENTION: DIAGNOSTIC USES THEREFOR
FILE REFERENCE: MNI-132CP3
CURRENT APPLICATION NUMBER: US/09/635,501
CURRENT FILING DATE: 2000-08-09
CURRENT FILING DATE: 2000-08-09
CURRENT FILING DATE: 2000-08-09
                                                                                                                                                                                         SEQ ID NO 29
LENGTH: 18
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 29, Application US/09635501 Patent No. 6884771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 13
LENGTH: 18
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 29.5%;
Best Local Similarity 54.5%;
Matches 6; Conservative
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Patent No. 657324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: GORDON, RICHARD K.
APPLICANT: MOORAD, DEBORAH R.
APPLICANT: DOCTOR, BHUPENDRA P.
APPLICANT: GARCIA, GREGORY E.
TITLE OF INVENTION: PREVINS AS SPECIFIC INHIBITORS AND THERAPEUTIC TITLE OF INVENTION: FOR BOTULINUM TOXIN B AND TETANUS NEUROTOXINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/134,446
PRIOR FILING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 38644-170531
CURRENT APPLICATION NUMBER: US/09/570,022
CURRENT FILING DATE: 2000-05-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
1 DPAKGMSPPGF 11 | :|| : || |
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                                                                  Similarity 6; Conserv
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                                                                                 29.5%;
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                                                                                                                                                                                                                                                                                                                                                                 08/989,299
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                                                             Score 31; DB 2; Louis Pred. No. 1.9e+02; 2; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 31; DB 2;
Pred. No. 1.9e+02
2; Mismatches
                                                                                                       Length 18;
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                                                                  Indels
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В

DISKGENNPGF 11

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APPLICANT: FOY, Teresa M.
APPLICANT: FOY, Teresa M.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210111.455C17
CURRENT APPLICATION NUMBER: US/10/007,700
CURRENT FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 469
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 463
LENGTH: 20
TYPE: PRT
ORGANISM: Homo sapiens
US-10-007-700-463
                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 22
US-08-475-955-12
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US-10-007-700-463
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                 Patent No. 6641813
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 69605
             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,955
FILING DATE: June 7, 1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                               APPLICANT: Harley, John
TITLE OF INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF
TITLE OF INVENTION: AUTOANTIBODIES
PRIOR APPLICATION DATA
                                                                                                                                                                                               STREET: 2800 O
CITY: Atlanta
STATE: GA
                                                                                                                                                                COUNTRY: USA
ZIP: 30309-3450
                                                                                                                                                                                                                              ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center, 1250 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
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Similarity 62.5%;
5; Conservative
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6641813
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PTSGMPPP 16
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Fanger, Neil
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Henderson, Robert A.
McNeill, Patricia D.
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Peckman, David W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vedvick, Thomas
Carter, Darrick
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Fanger, Gary R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Retter, Marc W.
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Pred. No. 2
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2.1e+02;
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Query Match
Best Local Similarity
Watches 5; Conserv:
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US-08-475-955-205
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              SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,955
FILING DATE: June 7, 1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
CLASSIFICATION DATA:
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
CLASSIFICATION DATA:
APPLICATION NUMBER: 07/648,205
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 205, Application US/08475955
Patent No. 6641813
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APPLICANT: Harley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (404)-873-879
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Harley, John TITLE OF INVENTION: METH TITLE OF INVENTION: AUTONUMBER OF SEQUENCES: 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMRF114CIP(2)DIV
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/648,205
FILLING DATE: January 31, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
NAME: DATE: THEORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center, 1250 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
FILING DATE: January 31, 1990 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/867,819 FILING DATE: April 13, 1992
                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Atlanta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Pabst, Patrea L. REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         METHODS AND REAGENTS FOR DIAGNOSIS OF AUTOANTIBODIES
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Pred. No. 4.6e+05;
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REGISTRATION NUMBER:

31,284

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APPLICANT: Harley,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 8 amino acids
                                                             TELEFAX: (404)-873-879
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                          CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
PRIOR APPLICATION DATA:
                                                                                                                        NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OWN
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                           APPLICATION NUMBER: 07/472,947 FILING DATE: January 31, 1990 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: 1i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF TITLE OF INVENTION: AUTOANTIBODIES NUMBER OF SEQUENCES: 218
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                            TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: Jur
CLASSIFICATION:
             STRANDEDNESS:
                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/475,955
FILING DATE: June 7, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Atlanta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: OMRF114CIP(2)DIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center, 1250 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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                                             8 amino acids
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(404)-873-8795
TD NO: 205:
o: single
linear
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83.3%;
                                                                                                                               31,284
ER: OMRF114CIP(2)DIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 30; DB 2;
Pred. No. 4.6e+05;
0; Mismatches 1
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Search completed: January 26, 2006, 08:07:04 Job time: 19.9598 secs

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US-07-867-819D-12
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                                                              Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                             SOFTWARE: P
                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Harley, John
TITLE OF INVENTION: Methods and Reagents for Diagnosis of Autoantibodies
FILE REFERENCE: OMRF 114 CIP (2)
CURRENT APPLICATION NUMBER: US/07/867,819D
CURRENT FILING DATE: 1992-04-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 28.6
Best Local Similarity 83.3
Matches 5; Conservative
                                                                                                                                          LENGTH: 8

TYPE: PRI
ORGANISM: homo sapien
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)...(8)
OTHER INFORMATION: Binding site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 12, Application US/07867819D Patent No. 6897287
                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 07/472,947
PRIOR FILING DATE: 1990-01-31
PRIOR APPLICATION NUMBER: 07/648,205
PRIOR FILING DATE: 1991-01-31
NUMBER OF SEQ ID NOS: 161
3 GMPPPG 8
                                5 GMSPPG 10
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Pred. No. 4.6e+05;
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  Published Applications AA Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

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US-10-147-255-67
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CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: US/09/292,225
PRIOR FILING DATE: 1999-04-15
PRIOR APPLICATION NUMBER: 60/098,909
PRIOR FILING DATE: 1998-09-02
PRIOR PELICATION NUMBER: 60/085,295
PRIOR APPLICATION NUMBER: 60/098,565
PRIOR APPLICATION NUMBER: 60/098,565
SEQ ID NO 13
LENGTH: 20
TYPE: PRT
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                                                                                                                                PRIOR APPLICATION NUMBER: 60/085,295
PRIOR FILING DATE: 1998-05-13
PRIOR APPLICATION NUMBER: 60/098,565
PRIOR FILING DATE: 1998-04-17
PRIOR APPLICATION NUMBER: 09/062,013
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SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: NOVEL DERMATOPHAGOIDES NUCLEIC ACID MOLECULES, TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: McCall, Catherine A. APPLICANT: Hunter, Shirley Wu APPLICANT: Weber, Eric R.
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PRIOR APPLICATION NUMBER: 09/062,013
PRIOR FILING DATE: 1998-04-17
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                                                                                        PRIOR FILING DATE: 1998-04-17
NUMBER OF SEQ ID NOS: 49
                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US/09/292,225
PRIOR FILING DATE: 1999-04-15
PRIOR APPLICATION NUMBER: 60/098,909
PRIOR FILING DATE: 1998-09-02
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CURRENT FILING DATE: 2002-08-13
                                                                     SOFTWARE: PatentIn Ver.
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Pred. No. 1.1e-08;
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; ORGANISM: Eschericia coli
FEATURE:
; NAME/KEY: MISC FEATURE
; OTHER INFORMATION: DGI5-20M-4-E4
US-10-280-066-302
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CURRENT FILING DATE: 2002-10-24
PRIOR APPLICATION NUMBER: 60/345,471
PRIOR FILING DATE: 2001-10-24
NUMBER OF SEQ ID NOS: 537
SOFTWARE: Patentin version 3.1
SEQ ID NO 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 302, Application US/10280066 Publication No. US20030180718A1
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Best Local Similarity
Matches 20; Conserv
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Best Local :
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APPLICANT: Dedova, Olga
APPLICANT: Blume, Arthur J.
APPLICANT: Prendergast, John
APPLICANT: Goldstein, Neil I.
TITLE OF INVENTION: TARGET SPECIFIC SCREENING AND ITS USE FOR IDENTIFYING TARGET BIND
FILE REFERENCE: 2598-4009US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Pillutla, Renuka
APPLICANT: Brissette, Renee
APPLICANT: Spruyt, Michael
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Local Similarity 53.3%;
les 8; Conservative
                ZIP: 30306-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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ADDRESSEE: Patrea L. Pabst
                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 38
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TITLE OF INVENTION: Diagnostics And Therapy Of Epstein-Barr
Virus In Autoimmune Disorders
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STATE: Georgia
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                                                                                                                                 COUNTRY: USA
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Pred. No. 3.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IDICALON NO.
GENERAL INFORMATION:
APPLICANT: Harley, John
TITLE OF INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF
AUTOANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: OMRF161
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404)873-8794
TELEPAX: (404)873-8795
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/781,296
FILING DATE: 13-JAN-1997
APPLICATION NUMBER: 60/019,053
FILING DATE: 16-MAY-1996
APPLICATION NUMBER: 08/160,604
FILING DATE: 30-NOV-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                               APPLICATION NUMBER: US/10/376,121A
FILING DATE: 27-Mar-2003
CLASSIFICATION: <UNknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                 ZIP: 30309-3400

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 218
CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE DESCRIPTION: SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 PAKGMSPP 9
                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Atlanta
STATE: GA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/10/012,756 FILING DATE: 24-Oct-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: Suite 2000, 1201 West Peachtree Street, N.E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Pabst, Patrea L. REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             32.4%;
75.0%;
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Pred. No. 1.7e+06;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                     Version #1.25
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US-10-376-121A-100
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; SEQUENCE DESCRIPTION: SEQ ID NO: 116:
US-10-376-121A-116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 100, Application US/10376121A
Publication No. US20030216544A1
GENERAL INFORMATION:
APPLICANT: Harley, John
TITLE OF INVENTION: METHODS AND REAGENTS
AUTOANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local S
                                                                                                                                                                                                    TELEFAX: (404)-817-8588 INFORMATION FOR SEQ ID NO: 100:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 116: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/376,121A

FILING DATE: 27-Mar-2003
SEQUENCE DESCRIPTION: SEQ ID NO: 100:
                                                                                                                                                                                                                                                          NAME: Pabst, Patrea L.

REGISTRATION NUMBER: 31,284

REFERENCE/DOCKET NUMBER: OMRF114CIP(2)DIV(2)

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 30309-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: Suite 2000,-1201 West Peachtree Street, N.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: OMRF114CIP(2)DIV(2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404)-817-8473
                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                            FEATURE:
                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 218
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                   NAME/KEY:
LOCATION:
                                                                                              TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Atlanta
STATE: GA
                                                                                                                                                                                                                                        TELEPHONE: (404)-817-8473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acids
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                                                                                                                                                           LENGTH: 9 amino acids
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                         Binding-site
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                                                                                                              US-09-953-510-67
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Sequence 67, Application US/09953510
Sequence 67, Application US/09953510
Patent No. US20020131975A1
GENERAL INFORMATION:
APPLICANT: Horwitz, Marcus A.
TITLE OF INVENTION: Abundant Extracellular
TITLE OF INVENTION: Products and Methods for Their Production and Use
                                    Query Match
Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (310) 277-1
INFORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/447,398
FILING DATE: 23-MAY-1995
APPLICATION NUMBER: US 08/289,667
FILING DATE: 12-AUG-1994
APPLICATION NUMBER: US 08/156,358
FILING DATE: 23-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: MacLean, Kurt A.
REGISTRATION NUMBER: 31.1
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                    ORIGINAL SOURCE:
ORGANISM: Mycobacterium tuberculosis
                                                                                                                                                                                                                                              TRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR
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                                                                                                                                  SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                           FRAGMENT TYPE: internal
                                                                                                                                                                                                                                 ANTI-SENSE: NO
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 1 DPAKGMSP 8
                                      Similarity
5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/953,510
FILING DATE: 14-Sep-2001
CLASSIFICATION: <Unknown>
R APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                      LENGTH: 15 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 2029 Century Park East, Suite 3800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                       STRAIN: Erdman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Kurt A. MacLean
                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                  (310) 788-5000
310) 277-1297
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75.0%;
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                                      Score 33; DB 3; Pred. No. 5.4e+02; 2; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          112-272
                                                                                                                                   67:
                                                                             Length 15;
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                                          Indels
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RESULT 9
US-10-147-255-67
US-10-147-255-67
Sequence 67, Application US/10147255
Publication No. US20030152584A1
GENERAL INFORWATION:
                                                                                                                                                                                                                                                                                         STRAIN: Erdman
;
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-953-413-67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-953-413-67; Sequence 67, Application US/09953413; Publication No. US20040018209A1
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Matches 5; Conserv
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APPLICANT: Horwitz,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/447,398
FILING DATE: 23-MAY-1995
APPLICATION NUMBER: US 08/289,667
FILING DATE: 12-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Kurt A. MacLean
STREET: 2029 Century Park East, Suite 3800
                                                                                                                                                                                                                                                                                                                                                                              FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/953,413
                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                                                  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                  DPSQGMGP 11
                                                                                                                                                                                    DPAKGMSP 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: MacLean, Kurt A. REGISTRATION NUMBER: 31,118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/156,358 FILING DATE: 23-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Mycobacterium tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (310) 788-5000
TELEFAX: (310) 277-1297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 14-Sep-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                   . peptide
                                                                                                                                                                                                                                             31.4%;
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Abundant Extracellular
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                                                                                                                                                                                                                                             Score 33; DB 3;
Pred. No. 5.4e+02;
                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     112-272
                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                               Length 15;
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US-10-695-155-67
Sequence 67. Application US/10695155
Publication No. US20040228873A1
GENERAL INFORMATION:
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                                                                                                                                                                                                        RESULT 10
                                                                                                                                                                                                                                                                                                                                                   Matches
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APPLICANT: HORWITZ, MARCUS A.
APPLICANT: HARTH, GUNTER
APPLICANT: LEE, BAI-YU
TITLE OF INVENTION: ABUNDANT EXTRACELLULAR PRODUCTS AND METHODS FOR THEIR PRODUCTION
TITLE OF INVENTION: AND USE
FILE REFERENCE: 51326-00004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (310) 788-5000
TELEFAX: (310) 277-1297
INFORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/226,539A FILING DATE: «Ubknown» APPLICATION NUMBER: US 08/447,398 FILING DATE: 23-MAY-1995 APPLICATION NUMBER: US 08/289,667 FILING DATE: 12-AUG-1994 APPLICATION NUMBER: US 08/156,358 FILING DATE: 23-NOV-1993 ATTORNEY,ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: Mycobacterium tuberculosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Horwitz, Marcus A.
TITLE OF INVENTION: Abundant Extracellular
Products and Methods for Their Production and Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Kurt A. MacLean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER
                                                                                                                                                                                                                                                                                                         1 DPAKGMSP 8
                                                                                                                                                                                                                                                                                                                                                 Similarity
5; Conser
                                                                                                                                                                                                                                                                   DPSQGMGP 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: <Unknown>
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REGISTRATION NUMBER: 31,118
REFERENCE/DOCKET NUMBER: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/10/147,255 FILING DATE: 15-May-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH: 15 amino acids
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Pred. No. 5.4e+02;
2; Mismatches 1;
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US-09-907-969-397
; Sequence 37, Application US/09907969
; Publication No. US20030091580A1
; GENERAL INFORMATION:
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Best Local Similarity (
                                                                                 RESULT 12
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US-09-884-441-397
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APPLICANT: Algate, Paul A.
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C7
                                                                                                                                                                                                                                                                                                                            SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 397
LENGTH: 20
TYPE: PRT
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Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/884,441
CURRENT FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 489
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PRIOR APPLICATION NUMBER: 08/156,358
PRIOR FILING DATE: 1993-11-23
PRIOR APPLICATION NUMBER: 08/545,926
PRIOR FILING DATE: 1995-10-20
NUMBER OF SEQ ID NOS: 167
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CURRENT FILING DATE: 2003-10-27
PRIOR APPLICATION NUMBER: 08/652,842
PRIOR FILING DATE: 1996-03-23
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PRIOR FILING DATE: 1995-12-06
PRIOR APPLICATION NUMBER: 08/551,149
PRIOR FILING DATE: 1995-10-31
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TYPE: PRT
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Pred. No. 7.4e+02;
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Pred. No. 5.4e+02;
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APPLICANT: Mitcham, Jennifer L. APPLICANT: King, Gordon E. APPLICANT: Algate, Paul A.

APPLICANT:

King, Gordon E. Algate, Paul A. Fling, Steven P. Retter, Marc W. Fanger, Gary Richard

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APPLICANT: Albone, Earl
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C8
CURRENT APPLICATION NUMBER: US/09/907,969
CURRENT FILING DATE: 2001-07-17
NUMBER OF SEQ ID NOS: 596
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 397
LENGTH: 20
TYPE: PAT
ORGANISM: Homo sapiens
US-09-907-969-397
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CURRENT FILING DATE: 2001-04-04
NUMBER OF SEQ ID NOS: 461
SOFTWARE: FBSETSEQ for Windows Version 3.0
SEQ ID NO 397
LENGTH: 20
TYPE: PRT
ORGANISM: Homo Bapiens
US-09-827-271-397
Sequence 397, Application US/10198053

Publication No. US20030124140A1

GENERAL INFORMATION:
APPLICANT: Bangur, Chaitanya S.
APPLICANT: Retter, Marc W.
APPLICANT: Hanger, Gary R.
APPLICANT: Hill, Paul
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C9
CURRENT APPLICATION NUMBER: US/10/198,053
CURRENT FILING DATE: 2002-07-17
NUMBER OF SEG ID NOS: 624
SOFTWARE: FastSEQ for Windows Version 4.0
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US-09-827-271-397
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APPLICANT: Fanger, Gary R.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.46266
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Local Similarity 62.5%;
hes 5; Conservative
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o. US20030165504A1
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Pred. No. 7.4e+02;
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Pred. No. 7.4e+02;
3; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/10/198,053
; CURRENT FILING DATE: 2002-07-17
; NUMBER OF SEQ ID NOS: 624
; SOPTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 603
; LENGTH: 20
; TYPE: PRT
; ORGANISN: Homo sapiens
US-10-198-053-603
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Best Local Similarity
Watches 5; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                              US-10-860-790-397; Sequence 397, Application No. US200; Publication No. US200; GENERAL INFORMATION:
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                                                                                                                       US-10-860-790-397
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                                                                                                                                                   TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C11
CURRENT APPLICATION NUMBER: US/10/860,790
CURRENT FILING DATE: 2004-06-02
NUMBER OF SEQ ID NOS: 624
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 397
LENGTH: 20
TYPE: PRT
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Matches 5; Conserv
                                                 Best Local Similarity
Matches 5; Conserv
                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                           APPLICANT: Bangur, Chaitanya S.
APPLICANT: Retter, Marc W.
APPLICANT: Fanger, Gary R.
APPLICANT: Hill, Paul
                                                                                                                                     ORGANISM: Homo sapiens
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13 VGEEGVLS 20
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ilarity 62.5%;
Conservative
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                                                                     Score 33; DB 5;
Pred. No. 7.4e+02;
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Pred. No. 7.4e+02;
3; Mismatches 0
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Pred. No. 7.
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Query Match
Best Local Similarity
Marches 5; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-661-156-146
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US-10-860-790-603
           CURRENT APPLICATION NUMBER: US/10/661,156
CURRENT FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: US 10/382,082
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: PCT/US03/06731
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US 60/440,411
PRIOR APPLICATION NUMBER: US 60/440,411
PRIOR APPLICATION NUMBER: US 60/360,851
PRIOR FILING DATE: 2002-03-01
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US-10-860-790-603
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SOFTWARE: FRSTSEQ for Windows Version 4.0
SEQ ID NO 603
LENGTH: 20
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                         APPLICANT: Von Wronski, Mathew A.
TITLE OF INVENTION: KDR and VEGF/KDR Binding Peptides and
TITLE OF INVENTION: Their Use in Diagnosis and Therapy
FILE REFERENCE: D0617.70012US00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 210121.462C11
CURRENT APPLICATION NUMBER: US/10/860,790
CURRENT FILING DATE: 2004-06-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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Linder, Karen E.
Marinelli, Edmund R.
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                                                                                                                                                                                                                                                                                                                                       Shrivastava, Ajay
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5. US20050031634A1
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for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Christophe
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Pred. No. 7.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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US-10-328-916-58
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LENGTH: 15

TYPE:.PRT

ORGANISM: Artificial Sequence
FEATURE:
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LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Publication No.
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Best Local Similarity
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Best Local Similarity
PRIOR APPLICATION NUMBER: 60/200,065
PRIOR FILING DATE: 2000-04-27
NUMBER OF SEQ ID NOS: 26
                                                        APPLICANT: Van Heuit, Pam T
APPLICANT: Wu, John
TITLE OF INVENTION: DNA Encoding a No. US20020009455A1el PROST
FILE REFERENCE: 51831AUSM1
CURRENT APPLICATION NUMBER: US/09/838,785
CURRENT FILING DATE: 2001-04-20
CURRENT FILING DATE: 2001-04-20
                                                                                                                                                                                                                                   APPLICANT: Lau, Ted
APPLICANT: Lin, Rick
APPLICANT: Parkes, D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Blume, John E.

APPLICANT: Metabolex, Inc.
TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor TITLE OF INVENTION: Involved in Islet Cell Signaling FILE REFERENCE: 016325-007210US
CURRENT APPLICATION NUMBER: US/10/328,916
CURRENT FILING DATE: 2003-01-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/345,697
PRIOR FILING DATE: 2002-01-04
NUMBER OF SEQ ID NOS: 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Gregoire, Francine M.
APPLICANT: Johnson, Jeffrey D.
APPLICANT: Blume, John E.
APPLICANT: Metabolex, Inc.
                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence:C-terminal OTHER INFORMATION: peptide of human IC-GPCR for raising antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: Library Isolate
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                                                                                                                                          Parkes, Debbie
Parry, Gordon
Schneider, Douglas
Steinbrecher, Renate
Van Heuit, Pam T
Wu, John
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Pred. No. 7.4e+02;
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Pred. No. 7.6e+02;
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US-09-838-785-26
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SEQ ID NO 26
LENGTH: 16
                Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 901
SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 593
LENGTH: 16
TYPE: PRT
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Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/953,901
CURRENT FILING DATE: 2004-09-29
CURRENT FILING NATE: PCT/CA03/00465
PRIOR APPLICATION NUMBER: PCT/CA03/00465
PRIOR FILING DATE: 2003-04-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: VIRAG, CRISTINA
TITLE OF INVENTION: NOVEL PURIFIED POLYPEPTIDES FROM BACTERIA
FILE REFERENCE: IPT-207.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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TYPE: PRT
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ORGANISM: Staphylococcus aureus
                                                                                                                                                                                                                                                                                                             R APPLICATION NUMBER: 60/369,831
FILING DATE: 2002-04-04
RAPPLICATION NUMBER: 60/369,819
FILING DATE: 2002-04-04
APPLICATION NUMBER: 60/369,826
                                                                                                                       FILING DATE: 2002-04-09
                                                                                                                                     APPLICATION NUMBER: 60/371,180 FILING DATE: 2002-04-09 APPLICATION NUMBER: 60/371,008
                                                                                                                                                                                           APPLICATION NUMBER: 60/371,014 FILING DATE: 2002-04-09
                                                                                                                                                                                                                            APPLICATION NUMBER: 60/370,681 FILING DATE: 2002-04-08
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APPLICATION NUMBER: 60/370,852
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VIRAG, CRISTINA
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ALAM, MUHAMMAD ZAHOOR
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MCDONALD, MERRY-LYNN
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HOUSTON, SIMON
KANAGARAJAH, DHUSHY
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BUZADZIJA, KRIS
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b. US20050181464A1
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Pred. No. 8.2e+02
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                                                                                                     See File Wrapper or PALM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 16;
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APPLICANT: Von Wronski, Mathew A.
TITLE OF INVENTION: KDR and VEGF/KDR Binding Peptides and
TITLE OF INVENTION: Their use in Diagnosis and Therapy
FILE REFERENCE: D0617.70012US00
CURRENT APPLICATION NUMBER: US/10/661,156
CURRENT FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: US 10/382,082
PRIOR APPLICATION NUMBER: US 10/382,082
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: PCT/US03/06731
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US 60/440,411
PRIOR FILING DATE: 2003-01-15
PRIOR APPLICATION NUMBER: US 60/440,411
PRIOR FILING DATE: 2003-01-15
PRIOR APPLICATION NUMBER: US 60/360,851
PRIOR FILING DATE: 2003-03-01
PRIOR FILING DATE: 2003-03-01
RESULT 23
US-10-307-956-25
; Sequence 25, Application US/10307956
; Publication No. US20030119072A1
; GENERAL INFORMATION:
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Best Local Similarity
"atches 7; Conserv
                                                                                                                        밁
                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Library Isolate US-10-661-156-170
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US-10-661-156-170
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                                                                                                                                                             Matches
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Best Local (
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                                                                                                                                                                                                                                                                                              LENGTH: 20
TYPE: PRT
ORGANISM: Artificial Sequence
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Linder, Karen E.
Marinelli, Edmund R.
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Bussat, Philippe
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Pochon, Sibylle
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o. US20050100963A1
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Pred. No. 1e+03;
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8.2e+02;
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APPLICANT: Hoekstra, Merl F.

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RESULT 25
US-10-950-163-8
Sequence 8, Application US/10950163
Publication No. US20050152911A1
GENERAL INFORMATION:
APPLICANT: Hardy, Michele
TITLE OF INVENTION: NOROVIRUS MONOCLONAL ANTIBODIES AND PEPTIDES
FILE REFERENCE: 33576/US/2
CURRENT APPLICATION UNMBER: US/10/950,163
CURRENT FILING DATE: 2004-09-24
PRIOR APPLICATION NUMBER: US 60/508,262
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US-10-211-088-356
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; TYPE: PRT
; ORGANISM: Homo sapien
US-10-307-956-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NO. US20030104479A1el Fusion Proteins And Assays For Molecular Bi FILE REFERENCE: 01-1022-US CURRENT APPLICATION NUMBER: US/10/211,088 CURRENT FILING DATE: 2002-10-15 PRIOR APPLICATION NUMBER: 60/309,395 PRIOR FILING DATE: 2001-08-01 PRIOR APPLICATION NUMBER: 60/341,589 PRIOR FILING DATE: 2001-08-01 PRIOR FILING DATE: 2001-08-01 PRIOR FILING DATE: 2001-08-01 Seq ID NOS: 366 SOFTMARE: Patentin version 3.1 SEQ ID NO 356
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Best Local Similarity 38.5%;
Matches 5; Conservative "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/307,956
CURRENT FILING DATE: 2002-12-02
PRIOR APPLICATION NUMBER: US/09/385,918
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEO ID NOS: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bright, Gary R. APPLICANT: Premkumar, D. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial sequence FEATURE:
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PGKGVKSPG 10
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Chen, Yih-Tai
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; PRIOR FILING DATE: 2003-09-24
; NUMBER OF SEQ ID NOS: 159
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 8
; LENGTH: 17
; TYPE: PRT
; ORGANISM: Grimsby virus
US-10-950-163-8

Query Match
Best Local Similarity 71.4%; Pred. No. 1.2e+03;
Matches 5; Conservative 2; Mismatches 0; Indels 0;
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2. PAKGMSP 8
Db 7 PAEGMSP 8
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Search completed: January 26, 2006, 08:38:40
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and is derived by \varepsilon
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Maximum
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/cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep: *

/cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep: *

/cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep: *

/cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep: *

/cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep: *

/cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep: *

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US-10-939-890-170
US-10-623-155-541
US-10-623-155-524
US-10-623-155-278
US-10-623-155-278
US-11-94-142-16
US-11-94-155-3003
US-11-158-735-60
US-11-158-61-78
US-11-154-61-78
US-11-154-61-78
US-11-054-515-2783
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                   Sequence
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                 146, App
160, App
541, App
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205, Appl
205, Appl
207, Appl
207
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29
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     US-11-147-994-9
US-10-939-890-13
US-10-939-890-13
US-10-839-584-240
US-10-842-266-26
US-10-503-575-131
US-10-997-2011A-31
US-10-623-155-545
US-10-623-155-545
US-10-623-155-253
US-10-623-155-254
US-10-623-155-265
US-11-045-024-1467
US-11-045-024-161
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 8 9, Appli
193, App
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240, Appp
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251, Appp
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261, Appp
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US-10-939-890-170
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US-10-939-890-146
                                                                 Sequence 170, Application US/10939890 Publication No. US20050250700A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 146
LENGTH: 20
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Best Local (
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                APPLICANT: Sato, Aaron K. APPLICANT: Sexton, Danie APPLICANT: Dransfield, D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2002-03-01
NUMBER OF SEQ ID NOS: 883
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 10/382,082
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: PCT/US03/06731
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US 60/440,411
PRIOR FILING DATE: 2003-01-15
PRIOR APPLICATION NUMBER: US 60/360,851
PRIOR FILING DATE: 2003-01-15
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APPLICANT: Swenson, Rolf E.
APPLICANT: Von Wronski, Mathew A.
TITLE OF INVENTION: KDR AND VEGF/KDR BINDING PEPTIDES
FILE REFERENCE: D0617.70014US00
 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 10/661,156 PRIOR FILING DATE: 2003-09-11
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7; Conserva
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Linder, Karen E.
Marinelli, Edmund R.
Manjappan, Palaniappa
Nunn, Adrian D.
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Dransfield, Daniel T.
Ladner, Robert C.
Sexton, Daniel J.
Dransfield, Daniel T.
Ladner, Robert C.
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Ramalingam, Kondareddiar
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                                                                                                                                                                                                                                                                        Conservative
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US-11-033-039-1324
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Pred. No. 19;
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Sequence 1324, Ap
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US-10-623-155-463
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                                                                         US-10-623-155-463
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SEQ ID NO 170
LENGTH: 20
                                                                                                                        CURRENT FILING DATE: 2003-07-17
NUMBER OF SEQ ID NOS: 560
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 463
LENGTH: 20
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Query Match
Best Local Similarity
Matches 5; Conserv
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APPLICANT:
APPLICANT:
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APPLICANT: Peckham, David W.
APPLICANT: Peckham, David W.
APPLICANT: Retter, Marc W.
APPLICANT: Fanger, Gary R.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C20
CURRENT APPLICATION NUMBER: US/10/623,155
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PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: US 10/382,082
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: PCT/US03/06731
PRIOR FILING DATE: 2003-03-03
PRIOR FILING DATE: 2003-03-03
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TITLE OF INVENTION: KOR AND VEGF/KDR BINDING PEPTIDES
FILE REFERENCE: D0617.70014US00
CURRENT APPLICATION NUMBER: US/10/939,890
CURRENT FILING DATE: 2004-09-13
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PRIOR FILING DATE: 2003-01-15
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                                                                                        TYPE: PRT ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30.5%;
Local Similarity 50.0%;
les 6; Conservation
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Linder, Karen E.
Marinelli, Edmund R.
Nanjappan, Palaniappa
Nunn, Adrian D.
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o. US20050261166A1
29.5%; ilarity 62.5%; Conservative
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; Score 31; DB;
; Pred. No. 39;
0; Mismatches
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Pred. No. 27;
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RESULT 6
US-10-623-155-524
/ Sequence 524, Application US/10623155
/ Publication No. US20050261166A1
/ GENERAL INFORMATION:
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SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 413
LENGTH: 20
TYPE: PRT
ORGANISM: Homo sapiens
US-10-623-155-413
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US-10-623-155-541
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US-10-623-155-541
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APPLICANT: Retter, Marc W.
APPLICANT: Franger, Gary R.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C20
CURRENT APPLICATION NUMBER: US/10/623,155
CURRENT FILING DATE: 2003-07-17
NUMBER OF SEQ ID NOS: 560
SOFTWARE: FABSEQ for Windows Version 4.0
SEQ ID NO 541
LENGTH: 20
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Best Local Similarity
Matches 5; Conserv
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Publication No. US20
GENERAL INFORMATION:
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Matches 5; Conserv
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APPLICANT: Peckham, David W.
APPLICANT: Retter, Marc W.
APPLICANT: Fanger, Gary R.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
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62.5%;
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Pred. No. 39;
0; Mismatches
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CURRENT FILING DATE: 2005-03-31
NUMBER OF SEQ ID NOS: 47
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 16
                                                                                                                                                                              Sequence 25, Application US/10889197 Publication No. US20050271689A1 GENERAL INFORMATION:
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Best Local Similarity 55.6%;
Matches 5; Conservative
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Publication No.
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Best Local Similarity
Matches 5; Conserv
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SEQ ID NO 524
LENGTH: 20
TYPE: PRT
APPLICANT: HUANG, CHUN-MING
APPLICANT: ZHANG, JIANFENG
APPLICANT: ZHANG, JIANFENG
APPLICANT: TANG, DE-CHU
TITLE OF INVENTION: NOVEL TARGETS AND COMPOSITIONS FOR USE IN
TITLE OF INVENTION: DECONTAMINATION, IMMUNOPROPHYLAXIS, AND POST-EXPOSURE
TITLE OF INVENTION: THERAPY AGAINST ANTHRAX
FILE REFERENCE: 858610-2006.1
CURRENT APPLICATION NUMBER: US/10/889,197
CURRENT FILING DATE: 2004-07-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Hagedorn, Peter H.
TITLE OF INVENTION: ANTIGEN ARRAY AND DIAGNOSTIC USES THEREOF
FILE REFERENCE: 29462
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APPLICANT:
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CURRENT FILING DATE: 2003-07-17
NUMBER OF SEQ ID NOS: 560
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APPLICANT: Peckham, David W.
APPLICANT: Reteer, Marc W.
APPLICANT: Fanger, Gary R.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 20
TYPE: PRT
GREANISM: Artificial sequence
FEATURE:
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Domany, Eytan
Elizur, Gad
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to. US20050260770A1
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Pred. No. 79;
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Pred. No. 79;
                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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US-10-939-890-203
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LENGTH: 12
                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 203
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                                             Matches
                                                                            Query Match
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PRIOR FILING DATE: 2003-07-11
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin Ver. 3.2
                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 10/661,156
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: US 10/382,082
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: PCT/US03/06731
PRIOR FILING DATE: 2003-03-03
PRIOR FILING DATE: 2003-03-03
                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: D0617.70014US00
CURRENT APPLICATION NUMBER: US/10/939,890
CURRENT FILING DATE: 2004-09-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                        LENGTH: 15
TYPE: PRT
                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 883
                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/360,851
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TYPE: PRT
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                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
                                             Local Similarity 71.4
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INVENTION: KDR AND VEGF/KDR BINDING PEPTIDES
2 PAKGMSP 8
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Similarity 55.6%;
5; Conservative
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Bussat, Philippe
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Ladner, Robert C.
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Nanjappan, Palaniappa
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Linder, Karen E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sibylle
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                                                             26.7%; Score 28; DB 71.4%; Pred. No. 81;
                                                0; Mismatches
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Pred. No. 63;
                                                                            DB 6; Length 15;
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 SOFTWARE: Pa
SEQ ID NO 212
                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 16
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Query Match
Best Local Similarity
Whiches 6; Conserve
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; ORGANISM: Homo sapiens
US-11-054-515-3003
                FILE REFERENCE: 10447-025

CURRENT APPLICATION NUMBER: US/10/893,584

CURRENT FILING DATE: 2004-07-19

PRIOR APPLICATION NUMBER: US 09/551,151

PRIOR FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: US 09/403,752

PRIOR FILING DATE: 1999-10-29

PRIOR FILING DATE: 1999-10-29

PRIOR APPLICATION NUMBER: US 10/089,058

PRIOR APPLICATION NUMBER: US 10/089,058

PRIOR APPLICATION NUMBER: US 10/089,058

PRIOR PILING DATE: 2000-10-04

NUMBER OF SEQ ID NOS: 274
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SEQ ID NO 3003
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
                                                                                                                                                                                                                                                                                       APPLICANT: Borgford, Thor
APPLICANT: Braun, Curtis
APPLICANT: Purac, Admir
APPLICANT: Stoll, Dominik
TITLE OF INVENTION: Ricin-Like Toxin Variants for Treatment of Cancer,
TITLE OF INVENTION: Viral or Parasitic Infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2004-06-18 PRIOR APPLICATION NUMBER: 10/293,418
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-10-17
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PRIOR APPLICATION NUMBER: 60/240,816
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FILING DATE: 2001-05-25
APPLICATION NUMBER: 60/277,379
FILING DATE: 2001-03-21
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PatentIn Ver.
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Best Local Similarity
Thes 5; Conserve
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: Sequence 78, Application US/11145861

: Publication No. US20060014138A1
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/ LENGTH: 20
/ TYPE: PRT
/ ORGANISM: E. coli
US-10-893-584-212

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US-11-058-735-60
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                                                                                                                 ; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-145-861-78
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; ORGANISM: Mus musculus
US-11-058-735-60
                                                                                                                                                  NUMBER OF SEQ ID NOS: 464
SOFTWARE: PatentIn version 3.3
SEQ ID NO 78
LENGTH: 11
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CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,748
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 80
SOFTWARE: PATENTIN Ver. 3.3
SEQ ID NO 60
LENGTH: 21
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Best Local Similarity
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                                                                                                                                                                                                                            APPLICANT: Chinnaiyan, Arul
APPLICANT: Wang, Xiaoju
TITLE OF INVENTION: Phage Microarray Profiling
FILE REFERENCE: UM-09899
CURRENT APPLICATION NUMBER: US/11/145,861
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3 FLEGEEQIL 11
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                                                                     25.7%;
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; Pred. No. 1.2e.
1; Mismatches
                                                       Score 27; DB Pred. No. 81; 2; Mismatches
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Pred. No. 1.1e+02;
"""matches 5;
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                                                                                                                                                       ; OTHER INFORMATION: PEPGEN P15 (TM) SYNTHETIC PEPTIDE US-10-516-676-1
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US-11-143-984A-164
                                                                                                                                                                                                                                                                                                                 SEQ ID NO 1
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                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/516,676
CURRENT FILING DATE: 2004-12-03
PRIOR APPLICATION NUMBER: 10/176,401
                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2002-06-21
NUMBER OF SEQ ID NOS: 1
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Philippart, Pierre
APPLICANT: Brasseur, Michele
TITLE OF INVENTION: SEALANT OR TISSUE GENERATING PRODUCT
FILE REFERENCE: 232037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/295,848
PRIOR FILLING DATE: 2001-06-05
PRIOR APPLICATION NUMBER: US 60/300,465
PRIOR FILING DATE: 2001-06-25
NUMBER OF SEQ ID NOS: 208
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PRIOR FILLING DATE: 2000-12-20
PRIOR APPLICATION NUMBER: US 60/280,186
PRIOR FILLING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: US 60/287,735
PRIOR FILLING DATE: 2001-05-01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING NOVEL HUMAN PHOSPHATASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 14
TYPE: PRT
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                                                                                                                                                                                            FEATURE:
OTHER INFORMATION: Synthesized
FEATURE:
                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                             LENGTH: 15
                                                                                            y Match 25.7%;
Local Similarity 40.0%;
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                                  5 GMSPPGFIVGEEGVL 19
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GTPGPQGIAGQRGVV 15
                                                                           Conservative
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                                                                                              Score 27; DB 6;
Pred. No. 1.2e+02;
                                                                           Mismatches
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2;
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RESULT 16 US-10-532-426-2

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Publication No. US20060009516A1
GENERAL INFORMATION:
APPLICANT: ROSENberg, Lawrence
TITLE OF INVENTION: USE OF INGAP FOR REVERSING
FILE REFERENCE: 1912-0308PUS1
CURRENT APPLICATION UNMBER: US/10/532,426
CURRENT FILING DATE: 2005-04-22
PRIOR APPLICATION UNMBER: US 60/420,677
                                                                                                                                                                                                       Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 2153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2153, Application US/11054515
Publication No. US20050255532A1
GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 2
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
                                                         Matches
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Best Local (
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
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PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                       ENGTH:
                                                                           Local
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/276,248
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/340,817
FILING DATE: 2001-12-19
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                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/240,816
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                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/277,379
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1 DPAKGMSPPGF 11
                                                   25.7%;
Similarity 54.5%;
6; Conservation
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Similarity 50.0%;
5; Conservative
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                                                   Score 27; DB /; LC
Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
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US-11-054-515-2783
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US-11-054-515-2780
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US-11-054-515-2780
                                                                                          PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/31,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
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PRIOR FILLING DATE: 2004-06-18
PRIOR PELLORATION NUMBER: 10/293,418
PRIOR FILLING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILLING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILLING DATE: 2001-12-19
PRIOR FILLING DATE: 2001-12-19
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SEQ ID NO 2780
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Best Local Similarity
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                       PRIOR FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 09/880,748
PRIOR FILING DATE: 2001-06-15
                                                                                                                                                                                                                                                                                                 APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
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CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
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PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
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APPLICATION NUMBER: 60/293,499
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Pred. No. 1.2e+02
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US-11-054-515-2783
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                                                                                                                             Sequence 598, Application US/10485788A Publication No. US20050282743A1
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SEQ ID NO 2783
                                                                                                        GENERAL INFORMATION:
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Best Local Similarity
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CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
                                               APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
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TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY PHASE TRANSITION
FILE REFERENCE: 4176-101 CIP
                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/240,816 PRIOR FILING DATE: 2000-10-17
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PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: REPEAT
LOCATION: (1)..(20)
OTHER INFORMATION: ELP1 [KIV2F1-4]
                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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CANT: Carrick, Deanna Marie
CANT: Arbor Vita Corporation
OF INVENTION: Molecular Interactions in Cells
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Pred. No. 1.6e+02;
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Pred. No. 1.2e+02
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US-11-178-269-27; Sequence 27, Application US/11178269; Publication No. US20050272661A1

GENERAL INFORMATION:

APPLICANT: Richelson, APPLICANT: Cusack, B APPLICANT: Pang, Yua

Cusack, Bernadette Marie Pang, Yuan-Ping McCormick, Daniel J.

Elliott

Faug, Abdul

RESULT 23

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                                                                                                                                                 ; OTHER INFORMATION: Small proteinaceous motif US-11-147-994-10
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-485-788A-598
                                                                      Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                 SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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Publication No.
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                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: CARPENTER, Kenneth W.
APPLICANT: TURNELL, William G.
APPLICANT: DeFIFE, Kristin M.
APPLICANT: GRAKO, Kathryn A.
TITLE OF INVENTION: BIOACTIVE STENTS FOR TYPE II DIABETICS AND METHODS FOR USE
TITLE OF INVENTION: THEREOF
FILE REFERENCE: MEDIV2030-3
CURRENT APPLICATION NUMBER: US/11/147,994
CURRENT FILING DATE: 2005-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 11/098,891
PRIOR FILING DATE: 2004-04-04
PRIOR APPLICATION NUMBER: US 60/559,937
PRIOR FILING DATE: 2004-04-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Medivas, APPLICANT: CARPENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/485,788A CURRENT FILING DATE: 2004-02-03 PRIOR APPLICATION NUMBER: US 60/309,841 PRIOR FILING DATE: 2001-08-03
                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                 LENGTH:
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                         PPGF 11
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                                                                        Conservative
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                                                                      24.8%; Score 26; DB 7; L
100.0%; Pred. No. 5.8e+04;
tive 0; Mismatches 0;
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Pred. No. 1.9e+02;
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TITLE OF INVENTION: NEO-TRYPTOPHAN
FILE REFERENCE: 07039/141001
CURRENT APPLICATION NUMBER: US/11/178,269
CURRENT FILING DATE: 2005-07-08

Beth Marie

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; FEATURE:
; OTHER INFORMATION: Synthetic peptide
US-11-043-788-482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 1998-04-10
NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 27
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetically generated peptide
US-11-178-269-27
                                                      RESULT 25
US-11-147-994-8
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PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US/10/265,099
PRIOR APPLICATION NUMBER: US/09/755,638
PRIOR FILING DATE: 2001-01-05
PRIOR APPLICATION NUMBER: 09/289,693
PRIOR FILING DATE: 2001-01-18
PRIOR APPLICATION NUMBER: US 60/098,119
PRIOR APPLICATION NUMBER: US 60/098,119
PRIOR APPLICATION NUMBER: US 60/098,119
PRIOR FILING DATE: 1998-08-27
Sequence 8, Application US/11147994 Publication No. US20060013855A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 482, Application US/11043788
Publication No. US20060014166A1
GENERAL INFORMATION:
APPLICANT: Compugen Ltd
TITLE OF INVENTION: NOVEL NUCLEOTIDE AND AMINO ACID SEQUENCES,
TITLE OF INVENTION: THEREOF FOR DIAGNOSIS OF ENDOMETRIOSIS
FILE REFERENCE: 1847.1006
                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 4; Conserv
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CURRENT FILING DATE: 2005-01-27
NUMBER OF SEQ ID NOS: 506
SEQ ID NO 482
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                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
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Pred. No. 5.8e+04;
1; Mismatches 0
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Mismatches 0;
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SEQ ID NO 8
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial sequence
                                                                              Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                             APPLICANT: Medivas, LLC
APPLICANT: CARPENTER, Kenneth W.
APPLICANT: TURNELL, William G.
APPLICANT: DeFIFE, Kristin M.
APPLICANT: DeFIFE, Kristin M.
APPLICANT: GRAKO, Kathryn A.
TITLE OF INVENTION: BIOACTIVE STENTS FOR TYPE II DIABETICS AND METHODS FOR USE
TITLE OF INVENTION: THEREOF
FILE REFERENCE: MEDIV2030-3
CURRENT APPLICATION NUMBER: US/11/147,994
CURRENT APPLICATION NUMBER: US 11/098,891
PRIOR APPLICATION NUMBER: US 11/098,891
PRIOR FILING DATE: 2004-04-04
PRIOR FILING DATE: 2004-04-05
NUMBER: OF SEQ ID NOS: 11
CORTMADE: Defeating 13
                                                                                                                                                                                   FEATURE: OTHER INFORMATION: Small proteinaceous motif
                                         8 PPGF 11
                                                                                   Conservative
                                                                                                    24.8%;
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                                                                                                  Score 26; DB 7; Li
Pred. No. 5.8e+04;
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                                                                                   Gaps
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Search completed: January 26, 2006, 08:39:07 Job time : 3.7931 secs

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Minimum DB
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Maximum Match
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Perfect score:
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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  GenCore version 5.1.6 (c) 1993 - 2006 Compugen Ltd
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B60588
E60788
I65478
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lectin, galactose/
50K allergen - per
photosystem ' nucleolin - bovine
vitronectin - goat
phosphocarrier por
immunodeficiency v
chymotrypsin (EC 3
neurofilament heav histone H2B - huma
GTP-binding protei
20K protein 5503 translation elonga
alpha-conotoxin Ep malate dehydrogena hypothetical 1.5K insulin-like growt complement factor proteasome chain 7
T-cell receptor al
pev-tachykinin - p
sperm-activating p group III allergen hypothetical prote fihrinoneptide A fibrinopeptide A - phosphocarrier pro vitronectin - shee cytochrome c(EDH) proteoglycan II, lipoprotein AcfD sperm-activating p sperm-activating p c-Ki-ras - hamster myosin heavy thyroid hormone re pyruvate dehydroge hypothetical prote glutathione transf cytochrome P450 transgelin nitrophorin histone H2B (clone histone H2B - huma sperm-activating 28K protein 4412 histone H2B - hum malate dehydrogena protein QA300033 -50K allergen - per photosystem I chai nephritogenoside -1-aminocyclopropan VCAM-1 5'UTR bindi chlorophyll a/b-bi sperm-activating photosystem I photosystem proteoglycan 65K glutathione peroxi DNA-directed RNA p insulin-like growt temperature-labile hypothetical myosin light 43.2K bile stone human leukocyte hypothetical prote sperm-activating sperm-activating ribosomal protein ribosomal protein lysyl-bradykinin cell receptor chain prote chain Ω Ö

transitional endop

ALIGNMENTS

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A; Cross-references: UNIPROT: Q7M2N0; UNIPARC: UPI000017C4E3
                                                                                                 A; Molecule type: protein A; Residues: 1-10 < JAN>
                                                                                                                                                                          A; Title: Further characterization of the three polypeptide chains A; Reference number: S26506; MUID:83209648; PMID:6852033
                                                                                                                                                                                                                                                C;Species: Bos primigenius taurus (cattle)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C;Accession: $26506
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Neuron 4, 697-709, 1990
A;Title: Differential splicing generates a
A;Reference number: PH0082; MUID:90262720;
A;Accession: PH0082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neuroglian protein, short form - fruit fly (Drosophila sp.) (fragment)
C;Species: Drosophila sp.
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH0082
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A; Accession: F57789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             submitted to the Protein Sequence Database, A;Description: The proteins of gallbladder:
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R;Binette, J.P.; Binette,
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                                                                                                                                                        A;Reference number: S26506;
A;Accession: S26506
                                                                                                                                                                                                                Eur. J. Biochem. 133, 39-46, 1983
                                                                                                                                                                                                                                  R; Jander, R.; Rautenberg, J.; Glanville,
                                                                                                                                                                                                                                                                                                             collagen alpha 1(VI) chain - bovine (fragment)
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A; Residues: 1-17 < HOR>
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;Date: 23-Feb-1996 #sequence_revision 23-Feb-1996 #text_change 09-Jul-2004
                                                                                                                                    Status: preliminary
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   Query Match
                                                           Keywords: hydroxyproline
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Similarity 60.0%;
6; Conservative
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                                       site: hydroxyproline (Pro)
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Pred. No.
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Pred. No. 1.3e+02;
3; Mismatches 6
     Score
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     28;
                                         #status experimental
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                                                                                                                                                                        4-hydroxybenzoate 3-monooxygenase (EC 1.14.13.2) - N;Alternate names: p-hydroxybenzoate 3-hydroxylase C;Species: Comamonas testosteroni
                                                                                               R;Tsugita, A. submitted to JIPID, May 1996
                                                                                                                                                                                                                                      RESULT
PT0089
A; Molecule type: proce
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A; Accession: PT0089 A; Reference number: PT0089 A; Description: Purification and

protein

characterization of p-hydroxybenzoate 3-hydroxylase

trom

;Date: 10-May-1996 #sequence_revision;Accession: PT0089

19-Jul-1996

#text_change

09-Jul-2004

Comamonas testosteroni (fragment)

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A;Cross-references: UNIPARC:UPI000017C5B4
C;Comment: spermatozoa must undergo capacitation and the acrosome reaction to become cap
C;Keywords: glycoprotein; semen
                                                                                                                                                                                                                                                           acrosome stabilizing factor small chain - rabbit (fragment) C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_ch C;Accession: B60801 C;Accession: B60801 R;Wilson, W.L.; Oliphant, G. Biol. Reprod. 37, 159-169, 1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: S54712

R;Chaudhuri, S.; Messing, J.

Mol. Gen. Genet. 246, 707-715, 1995

A;Title: RFLP mapping of the maize dzrl locus, which regulates methionine-rich A;Reference number: S54712; MUID:95206245; PMID:7898438

A;Accession: S54712
                                              Query Match
Best Local Similarity
Thes 5; Conserve
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;Species: Zea mays (maize)
;Date: 23-Aug-1995 #sequence_revision 19-Oct-1995 #text_change 09-Jul-2004
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                             DPAKGMSPPG 10
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Pred. No. 3.4e+02;
3; Mismatches 4
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Pred. No. 4.7e+02;
Pred. No. 5;
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RESULT 9

S65433

R;Wijffelb, G.; Fitzgerald, C.; Gough, J.; Riding, G.; Elvin, C.; Kemp, D.;
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C;Superfamily: 4-hydroxybenzoate 3-monooxygenase
C;Keywords: FAD: homodimer. cviaccai....
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A; Accession: PS0210
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A;Experimental source: patient EV, IL-2R+ synovial T-cells
A;Note: sequence extracted from NCBI backbone (NCBIP:113265)
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-16 <SIO>
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A;Title: Limited heterogeneity of T cell receptor variable region gene usage in juvenil A;Reference number: A49048; MUID:92387250; PMID:1387614
A;Accession: C49048
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                                                                                                                                                                                                                                                                                            ;Cross-references: UNIPROT:Q7M281; UNIPARC:UPI000017B0F9;Experimental source: germ, bran, strain Nihonbare;Comment: molecular weight 28K, pI 7.3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BK protein 4209 - rice (strain Nihonbare) (fragment)
;Species: Oryza sativa (rice)
;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Species: Ĥomo sapiens (man)
;Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 30-May-1997
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 Fitzgerald, C.; Gough, J.; Riding,
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Pred. No. 5.2e+02;
3; Mismatches 1
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Pred. No. 5.
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ornitho-kinin - chicken
C;Species: Gallus gallus (chicken)
C;Date: 11-Dec-1992 #sequence_revision 11-Dec-1992 #text_change 16-Aug-2004
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B60246
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A43065
                                                                                                                                         A; Cross-references: UNIPROT: Q7LZ50; UNIPARC: UPI000017A4F8
                                                                                                                                                            A;Molecule type: protein
A;Residues: 1-9 <KIM>
                                                                                                                                                                                                                               R;Kimura, M.; Sueyoshi, T.; Morita, T.; Tanaka, K.; Iwanaga, S. Adv. Exp. Med. Biol. 247A, 359-367, 1989
A;Title: Ornitho-kininogen and ornitho-kinin: isolation, characterization and A;Reference number: A60246; MUID:90102072; PMID:2603803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: UNIPROT:Q7LZI7; UNIPARC:UPI000002CF4A
C;Keywords: bradykinin; hydroxyproline; skin
F;3/Modified site: hydroxyproline (Pro) #status experimen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Nakajima, T.; Yasuhara, T.; Erspamer, G.F.; Visser, J. Experientla 35, 1133, 1979
A;Title: Occurrence of Hyp(3)-bradykinin in methanol extracts A;Reference number: A43065; MUID:80024576; PMID:488255
A;Accession: A43065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hydroxyproline-3-bradykinin - frog (Heleophryne purcelli)
C;Species: Heleophryne purcelli
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: protein A; Residues: 1-9 < WIJ>
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A;Title: Cloning and characterisation of angiotensin-converting
A;Reference number: 865431; MUID:96215437; PMID:8647080
A;Accession: $65433
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A;Residues: 1-9 <NAK>
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peptide -

garden dagger wasp

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k;Conion, J.M.; Hicks, J.W.; Smith, D.D.
Endocrinology 126, 985-991, 1990
A;Title: Isolation and biological activity of a novel kinin ([Thr(6)]bradykinin)
A;Reference number: A60579; MUID:90126625; PMID:2298179
A;Accession: A60579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Yasuhara, T.; Mantel, P.; Nakajima, T.; Piek, T. Toxicon 25, 527-535, 1987
A;Title: Two kinins isolated from an extract of the venom A;Reference number: A94322; MUID:87293024; PMID:3617088
A;Accession: A26744
                                                                                                                                         A;Molecule type: protein
A;Residues: 1-9 <CON>
A;Residues: 1-9 <CON>
A;Cross-references: UNIPARC:UPI000012DF29
C;Comment: This peptide increases aortic blood flow but, unlike bradykinin in mammalian C;Keywords: plasma
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C;Species: Megascolia flavifrons (ga
C;Date: 08-Mar-1989 #sequence_revis:
C;Accession: A26744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Piek, T.; Hue, B.; Mantel, P.; Nakajima, T.; Pelhate, M.; Yasuhara, T. Comp. Biochem. Physiol. C 96, 157-162, 1990
A;Title: Threonine(6)-bradykinin in the venom of the wasp Colpa interrupta A;Reference number: A61057; MUID:91130217; PMID:1980872
A;Accession: A61057
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C;Species: Pseudemys scripta (slider)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 05-Oct-2004
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C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 05-Oct-2004
C;Accession: A61057
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A;Residues: 1-9 <PIE>
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A;Residues: 1-9 <YAS>
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;Date: 08-Mar_1989 #sequence_revision 08-Mar-1989 #text_change 05-Oct-2004
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RESULT 17
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S13279
Ile-Ser-bradykinin - human (fragment)
N,Alternate names: T-kinin
C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: $13279
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C;Species: Rana temporaria (common frog)
C;Date: 09-Sep-1994 #sequence_revision 0
C;Accession: A61363
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                                                                                                                                         C;Accession: S13279
R;Wunderer, G.; Walter, I.; Eschenbacher, B.; Lang, M.; Kellermann, J.; Kindermann, G. Biol. Chem. Hoppe-Seyler 371, 977-981, 1990
A;Title: Ile-Ser-bradykinin is an aberrant permeability factor in various human malignan. A;Reference number: S13279; MUID:91166748; PMID:2076202
A;Accession: S13279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chem. Pharm. Bull. 16, 769-770, 1968
A;Title: Occurrence of a new active peptide on smooth muscle
A;Reference number: A61358; MUID:68412013; PMID:5677638
A;Accession: A61358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bradykinin-like peptide I - Japanese pond frog C;Species: Rana nigromaculata (Japanese pond frog) C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 C;Accession: A61358
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A61358
                                                                 A;Cross-references: UNIPROT:Q7M4P1; C;Keywords: bradykinin
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A;Residues: 1-9 <NAK>
A;Cross-references: UNIPROT:Q7LZ54; UNIPARC:UPI000017A4F0
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RESULT 20
A61360
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A61365
vespakinin M - hornet (Vespa mandarinia) C;Specias: Vespa mandarinia C;Date: 09-Sep-1994 #sequence_revision 0 C;Accession: A61360
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F;11/Binding site: sulfate (Tyr) (covalent)
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A;Title: Pharmacological data on phyllokinin (bradykinyl-isoleucyl-tyrosine A;Reference number: A61365; MUID:67179312; PMID:5970899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        phyllokinin - Rohde's leaf frog
N;Alternate names: bradykinyl-isoleucyl-tyrosine O-sulfate
C;Species: Phyllomedusa rohdei (Rohde's leaf frog)
C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_c
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Toxicon 25, 527-535, 1997
A;Title: Two kinins isolated from an extract of the venom reservoirs of the solitary was A;Reference number: A94322; MUID:87293024; PMID:3617088
A;Accession: B26744
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A;Accession: A28609
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R;Nakajima, T.; Piek, T.; Yashuara, T.; Mantel, P.
Toxicon 26, 34, 1988
A;Title: Two kinins isolated from the venom of Megascolia
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N;Alternate names:
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;Alternate names: 6-Thr-bradykinin-Lys-Ala
;Aspecies: Megascolla flavifrons (garden dagger wasp)
;Date: 08-Mar-1989 #sequence_revision 08-Mar-1989 #text_change 05-Oct-2004
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                #sequence_revision 09-Sep-1994 #text_change 05-Oct-2004
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RESULT 21
A61359
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A;Molecule type: protein
A;Residues: 1-13 <YAS>
                                                                                                                                                                                                             R;Yasuhara, T.; Hira, M.; Nakajima, T.; Yanaihara, N.; Yanaihara, C.; Hashimoto, Chem. Pharm. Bull. 21, 1388-1391, 1973
A;Title: Active peptides on smooth muscle in the skin of Bombina orientalis Bould A;Reference number: A61361; MUID:73256822; PMID:4732297
                                                                                                                                                                                                                                                                                                   C; Accession:
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C;Accession: A61361
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C;Species: Bombina orientalis
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A;Title: Chemical investigation of the hornet (Vespa xanthoptera A;Reference number: A61359; MUID:87187853; PMID:264186
A;Accession: A61359
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C;Species: Vespa xanthoptera
C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 05-Oct-2004
C;Accession: A61359
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C;Keywords: hydroxyproline; venom
F;4/Modified site: 4-hydroxyproline (Pro) #status experimental
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A;Title: Vespakinin-M, a novel bradykinin analogue containing hydroxyproline, A;Reference number: A61360; MUID:77114342; PMID:1017116
A;Recession: A61360
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24.8%; Score 26; DB 2; Length 13; 100.0%; Pred. No. 6e+02; rive 0; Mismatches 0; Indels
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100.0%;
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PPGF 5

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bradykinin-like peptide III - Japanese pond frog
;Species: Rana nigromaculata (Japanese pond frog)
C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 05-Oct-2004
C;Accession: A61362
A;Cross-references: UNIPROT:P57672; UNIPARC:UPI0000138459
C;Superfamily: vespulakinin
C;Keywords: antihypertensive; bradykinin; glycoprotein; venom
F;1-17/Product: vespulakinin 1 #status experimental <MAT1>
F;3-17/Product: vespulakinin 2 #status experimental <MAT2>
F;3-17/Product: vespulakinin 2 #status experimental <MAT2>
F;3-17/Region: bradykinin-like
F;3,4/Binding site: carbohydrate (Thr) (covalent) #status expe
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A36279
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A;Title: On the third active peptide on smooth muscle in the skin of Rana nigromaculata A;Reference number: A61362; MUID:69117202; PMID:5751736
A;Accession: A61362
A;Status: preliminary
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Biochemistry 15, 61-64, 1976
A;Title: Vespulakinins: new carbohydrate-containing bradykinin derivatives.
A;Reference number: A61339; MUID:76114777; PMID:1247511
A;Accession: A61339
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N;Contains: vespulakinin 2
C;Species: Vespula maculifrons (eastern yellowjacket)
C;Date: 17-Jul-1994 #sequence_revision 07-Oct-1994 #text_change 09-Jul-2004
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C;Species: Lumbricus terrestris (common earthworm)
C;Date: 18-Jan-1991 #sequence_revision 18-Jan-1991 #text_change 09-Jul-2004
C;Accession: A36279
C;Accession: A36279
R;Jiang, X.C.; Inouchi, J.; Wang, D.; Halpern, M.
J. Biol. Chem. 265, 8736-8744, 1990
A;Title: Purification and characterization of a chemoattractant from electric shock-indu
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Query Match 24.8%; Score 26; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 8.1e+02; Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0; Qy 8 PPGF 11

Db 10 PPGF 13
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Search completed: January 26, 2006, 08:05:06

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1P9 HUMAN

Q7M4P9 HUMAN 1

Q7M4P9;

Q1-MAR-2004 (
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25-OCT-2004 (TrEMBLrel. 28, Last sequence
25-OCT-2004 (TrEMBLrel. 28, Last annotate
25-OCT-2004 (TrEMBLrel).
208 proteasome beta subunit (Fragment).
Vitis vinifera (Grape).
Eukaryota; Viridiplantae; Streptophyta;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Carvalho M.B., Caeiro A.S., Submitted (JUL-2004) to the EMBL; AY684130; AAU04834.1; GO; GO:0005829; C:cytosol; I
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01-MAR-2004 (TYEMBLrel. 26, Last sequence update)
01-MAR-2004 (TYEMBLrel. 26, Last annotation update)
Gallbladder stone matrix protein 2, 41K (Fragment).
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Submitted (FEB-1996) to the
PIR; F57789; F57789.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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      (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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1808 MW;
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EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                                                     Score 30; DB 2;
Pred. No. 1.9e+03;
1; Mismatches (
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eudicotyledons; core eudicots; rosids;
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8.3e+02;
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rimates; Catarrhini; Hominidae;
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RESULT 5
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DE Zein Zp22/6 px
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OC Eukaryota; Viz
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OC PACCAD clade;
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Best Local :
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NUCLEOTIDE SEQUENCE.
MEDLINE=22294456; PubMed=12407086;
Kida M., Ariga T., Shirakawa T., Oguchi H., Sakiyama Y.;
Kida M., Ariga T., Shirakawa T. form of amelogenesis impe
   Eukaryota; Viridiplantae; Streptophyta; En
Spermatophyta; Magnoliophyta; Liliopsida;
PACCAD clade; Panicoideae; Andropogoneae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NON TER
SEQUENCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence up
01-MAR-2004 (TrEMBLrel. 26, Last annotation
Collagen alpha i(VI) chain (Fragment).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                          zeın Zp22/6 protein.
Zea mays (Maize).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jander R., Rauterberg J., Glanville R.W.;
"Further characterization of the three polypeptide and human short-chain collagen (intima collagen).";
Eur. J. Biochem. 133:39-46(1983).
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NCBI_TaxID=9913;
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                                                                                                                                                                                                                                                                                          PRELIMINARY;
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965 MW;
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Pred. No. 2.3e+03;
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                              STRAIN=ATCC 4797;

Langenheim J.F., Ulrich R.L.;

Submitted (WAR-2002) to the EMBL/GenBank/DDBJ databases.

EMBL, AF496531, AAQ07223.1; -; Genomic_DNA.

GO; GO:0016974; F:histidine-tRNA ligase activity; IEA.

GO; GO:0016974; F:ligase activity; IEA.

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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Histidyl-renn synthetase (EC 6.1.21) (Fragment).
Lactobacillus delbrueckii (subsp. lactis).
Bacteria; Firmicutes; Lactobacillales; Lactobacilla
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MEDLINE=95206245; PubMed=7898438; DOI=10.1007/BF00290716; Chaudhuri S., Messing J.;

"RFLP mapping of the maize dzrl locus, which regulates me 10 kDa zein accumulation.";
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Gonzalez Perez M.P.,
Submitted (JUL-2002)
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Human T-lymphotropic virus 1.
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PIR; S54712; S54712.
SEQUENCE 15 AA; 1605 MW; 5DFB4
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AF529962; AAP87693.1;
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Query Match
Best Local S
Matches 4
Pfam; PFO
NON_TER
SEQUENCE
                 PIR; PT0089; PT0089.

HSSP; P20586; IIUT.
GO; GO:0018659; F:4-hydroxybenzoate
InterPro; IPR002938; Moxy_FAD_bindin
Pfam; PF01494; FAD_binding_3; 1.

NON_TER
21
21
                                                                                                                                                                                                                           01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
4-hydroxybenzoate 3-monooxygenase (EC 1.14.13.2) (Fragment).
Comamonas testosteroni, (Pseudomonas testosteroni).
                                                                                                                     Tsugita A.
Submitted
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Q7M1D8_COMTE PRELIMINARY;
                                                                                                                                                                                 Comamonadaceae;
NCBI_TaxID=285;
                                                                                                                                                                                                                          Bacteria; Proteobacteria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; U40388; AAA83007.1; -; Genomic_DNA. HSSP; P20586; 1K0I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wong C.M., Dilworth M.J., Glenn A.R.;
"4-Hydroxybenzoate hydroxylase (pobA) is
in Rhizobium leguminosarum bv. viciael.";
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Q79CF0;
                                                                                                                                                       PROTEIN SEQUENCE.
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Pfam; PF01494; FAD_binding_3;
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Rhizobiaceae; Rhizobium/Agrobacterium group; Rhizobium
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
4-hydroxybenzoate hydroxylase (EC 1.14.13.2) (Fra
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Pred. No.
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Pred. No. 4.7e+03;
   52178D0D46B76055
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Matches 3
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                                                                                                                     Matches
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Q97204 BRANA PRELIMINARY;
Q97204;
Q97206;
01-MAY-2000 (TrEMBLrel. 13, C:
01-WAY-2000 (TrEMBLrel. 21, L:
Chaperonin-60 LS3 fragment (F:
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Q7M281;
01-MAR-2004 (
                                                                                                                                                                                                                                                                             MEDLINE=94302168; PubMed=7913238; DOI=10.1104/pp.105.1.233; Cloney L.P., Bekkaoui D.R., Feist G.L., Lane W.S., Hemmingsen S.I. "Brassica napus plastid and mitochondrial chaperonin-60 proteins contain multiple distinct polypeptides."; Plant Physiol. 105:233-241(1994).
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SEQUENCE
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;

Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mitochondrion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brassica napus (Rape)
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1579 MW;
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Q9N195_BOVIN PRELIMINARY;
AC Q9N195;
DT 01-OCT-2000 (TrEMBLrel. 15,
DE PARVAIDMININ (Fragment).
OS BOS TAURUS (BOVINE).
OC EUKARYOTA; Metazoa; Chordat
OC Mammalia; Eutheria; Laurasi
OC PECORA; BOVIDAE; BOVINAE; BOVINAE;
ON MEDILINE=2159954; PubMed=11
RA MEDILINE=2159954; PubMed=11
RA ARIZA F., HARRISON B., Drin
RT "The assignment by linkage
RT 22 to bovine chromosome 5 a
RL Anim. Genet. 32:371-374(200
DR EMBL; AF217654; AAF26420:1;
FT NON TER 1
PT NON TER 1
SEQUENCE 19 AA; 2151 MW;
                                            98 SULT POSSESSION OF SERVICE 
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J. Immunol.
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                                                                     EMBL; M34542; AAA39607.1;
NON_TER 1 1
NON_TER 21 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus (Mouse)
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Mammalia; Eutheria; Laurasiatheria;
                                               SEQUENCE
                                                                                                                                                                                                                                                                                            TISSUE=L
                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
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Flavell R.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=21891563; PubMed=11894963;
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5; Conserv
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                                               21
                                                                                                                                              147:3180-3184(1991)
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Q95554, MOUSE PRELIMINARY; PRT; 21 AA.
Q95554;
Q95554;
Q95554;
Q95554;
Q95564;
Q95564;
Q1-FEB-1997 (TrEMBLrel. 02, Created)
Q1-FEB-1997 (TrEMBLrel. 02, Last sequence update)
Q1-QCT-2002 (TrEMBLrel. 22, Last annotation update)
Major histocompatibility complex antigen H-2Kbm29 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=21599054; PubMed=11736808; Ariza F., Harrison B., Drinkwater R.D. "The assignment by linkage mapping of 22 to bovine chromosome 5 and 17.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Euarchontoglires; Glires; Rodent Muridae; Murinae; Mus.
                                                                                                                                                            MEDLINE-92013151; PubMed=1717590;
Horton R.M., Loveland B.E., Parwani A., Pease L.R., Lindahl K.F.;
"Characterization of the spontaneous mutant H-2Kbm29 indicates th
gene conversion in H-2 occurs at a higher frequency than detected
                                                                                                                                                                                                                                                                                                                                                 "The DNA sequence of the H-2Kb gene: Evidence mechanism for the generation of polymorphism i
                                                                                                                                                                                                                                                                                                                                                                                                               R., Mellor A.,
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Last sequence up
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Pred. No.
                                      103EC45620595224 CRC64;
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BRK1_RANNI
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Matches 4
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TISSUE=Skin secretion;

MEDLINE=22217713; PubMed=12230583;

Chen T., Orr D.F., Bjourson A.J., McClean S.,

Rao P., Shaw C.;

"Novel bradykinins and their precursor cDNAs
bellied toad (Bombina variegata) skin.";

Eur. J. Biochem. 269:4693-4700(2002).
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Q7LZ54;
                                                                                                                                                                                                                                                                                                                              Bombina variegata (Yellow-bellied toad).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Barrachia; Anura; Archeobatrachia; Bombinatoridae; Bomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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Bradykinin; Direct p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rana nigromaculata (Japanese pond frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata,
Amphibia, Batrachia, Anura, Neobatrachia, Ranoidea,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2003 (Rel.
28-FEB-2003 (Rel.
13-SEP-2005 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
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10-MAY-2005 (Rel. 47, Last annotation updat
Bradykinin-like peptide I.
                                                                                                                                                                                                                                          PROTEIN SEQUENCE,
                                                                                                                                                                                                                                                                                                NCBI_TaxID=8348;
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Pred. No.
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                                                                                                                                                                                                                                          LOCATION,
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2.2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 smooth muscle
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                                                                                                                                            O'Rourke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 9;
                                                                             European yellow-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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Ranidae; Rana
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                                                                                                                                                Hirst D.
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RESULT 16
KNL3_CYPDO
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Best Local S
Matches 4
             GO; GO:00056
GO; GO:00457
GO; GO:00459
Bradykinin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KNL3 CYI
P83659;
                                                                                                                                                                                                                                                                                     MEDLINE-21203862; PubMed=11306139; DOI-10.1016/S0041-0101(00)00262-2; Konno K., Hisada M., Naoki H., Itagaki Y., Yasuhara T., Juliano M.A., Juliano L., Palma M.S., Yamane T., Nakajima T.; "Isolation and sequence determination of peptides in the venom of the spider wasp (Cyphononyx dorsalis) guided by matrix-assisted laser desorption/ionization time of flight (MALDI-TOF) mass spectrometry."; Toxicon 39:1257-1260(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2004 (Rel. 44, 05-JUL-2004 (Rel. 44, 10-MAY-2005 (Rel. 47,
                                                                                                    use
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                                                                                                                                              This
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                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Venom;
                                                                                                                                                                                                                                                                                                                                                                                                           SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=246266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Vespoidea; Pompilidae; Cyphononyx.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cyphononyx dorsalis (Spider wasp)
Eukaryota; Metazoa; Arthropoda; H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use as long as its content is in no way modified and this statement is no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ++
                                                                                                                                                                                                                                                                                                                                                                                                                      PROTEIN SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vasoactive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amphibian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               removed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         [Thr6]bradykinin
GO:0005615; C:extracellular space; IDA.
GO:0045776; P:negative regulation of blood pres
GO:0045987; P:positive regulation of smooth mus
dykinin; Direct protein sequencing; Vasoactive;
UENCE 9 AA; 1074 MW; 3393D771A9C86777 CRC64;
                                                                                                                                                                                     TISSUE SPECIFICITY: Expressed by the venom gland.
MASS SPECTROMETRY: MW=1074.58; METHOD=MALDI; RANGE=1-9;
NOTE=Ref.1.
                                                                                                                s Swiss-Prot entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                         SIMILARITY:
                                                                                                                                                                                                                 SUBCELLULAR LOCATION: Secreted TISSUE SPECIFICITY: Expressed 1
                                                                                                     as
                                                                                                                                                                                                                                                           FUNCTION: Produces in vitro muscle and constriction of i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expressed by the skin gla
SIMILARITY: Belongs to the bradykinin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FUNCTION: Produces in vitro relaxation of rat arterial smooth muscle and constriction of intestinal smooth muscle.
                                                                                                                                                                                                                                               similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CYPDO
                                                                                                  long as its content is in no way modified and this statement
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   defense peptide;
; Vasodilator.
9 AA; 1074 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                                                                                                                                                                      Belongs to the bradykinin family.
                                                                                                                                                                                                                                                                                                                                                                                                                       SUBCELLULAR LOCATION, TISSUE SPECIFICITY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Last annotation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score; Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
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                                                                                                                                                                                                                                                           o relaxation intestinal :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hexapoda; Insecta;
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hes 0;
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                                                                                                                                                                                                                                                          n of rat arterial smooth smooth muscle (By
  CRC64;
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                                           pressure;
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                Vasodilator
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                                            ISS.
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Query Match Best·Local S Matches 4

Similarity
4; Conserv

24.8%; Silarity 100.0%; Conservative 0;

Score 26; DB 1; L Pred. No. 2.2e+06; 0; Mismatches 0;

Length 9;

Indels

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Gaps

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PPGF

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RESULT 19
Q7LZJ8 RANTE FID Q7LZJ8;
AC Q7LZJ8;
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Matches 4
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Best Local (
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77 TRASC
PB4497
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750 CHICK

Q7LZ50 CHICK |

Q7LZ50;

Q7LZ50;

Q7LZ50;

Q7LZ50;
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PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; (
Archosauria; Aves; Neognathae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-2004 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endocrinology 126:985-991(1990).
Direct protein sequencing.
SEQUENCE 9 AA; 1074 MW; 3393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Trachemys scripta (Red-eared slider turtle) (Pseudemys scripta) Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom Testudines; Cryptodira; Testudinoidea; Emydidae; Trachemys.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chemical structure.";
Adv. Exp. Med. Biol. 247A:359-367(1989)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=90102072; PubMed=2603803;
Kimura M., Sueyoshi T., Morita T., Tar
"Ornitho-kininogen and ornitho-kinin:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROTEIN SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ornitho-kinin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conlon J.M., Hicks J.W., Smith D.D.; "Isolation and biological activity of ([Thr(6)]bradykinin) from the turtle,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROTEIN SEQUENCE.
PubMed=2298179;
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10-MAY-2005 (TrEMBLrel. 30, Last sequence update)
10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Exp. 1. B60246; B'
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AA; 10
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                                               PRELIMINARY;
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Pred. No. 2.2e+06;
0; Mismatches 0;
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; Galliformes; Phasianidae; Phasianinae;
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RESULT
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JLT 21 p PHYRO PRIKE PHYRO Q7LZ52; 05-JUL-2004 (05-JUL-2004 (05-JUL-2005 (Phyllokinin (Phyllomedusa

STANDARD;

PRT;

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(Rel. 44, Created)
(Rel. 44, Last sequence update)
(Rel. 47, Last annotation update)
(Bradykinyl-isoleucyl-tyrosine Oa rohdei (Rohde's leaf frog).

O-sulfate).

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RESULT 20
Q7LZI7 9NEOB
Q7LZI7 9NEOB
Q7LZI7;
ID Z7
AC Q7LZI7;
AC Elephryne Metazoa; Chordat.
AC Amphibia; Batrachia; Anura;
AC NACHITANIA;
AC Amphibia; Batrachia; Anura;
AC NACHITANIA;
AC AMPHIBIA;
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PIR; A61363; A61363.
REQUENCE 9 AA; 1060 MW;
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PIR; A43065; A43065.
SEQUENCE 9 AA; 1060 MW; 3393D775B9C86777 CRC64;
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Heleophryne purcelli (Cape ghost frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Heleophrynidae;
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
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"Pharmacological data on phyllokinin (bradykinyl-isoleucyl-tyrosine
"Pharmacol and bradykinyl-isoleucyl-tyrosine.";
Br. J. Pharmacol. 27:479-485(1966).
-I- SUBCELLULAR LOCATION: Secreted.
                   uee
                                                              This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                           Nakajima T., Piek T., Yashuara T., Nakajima T., Nakaj
                                                                                                                                                                                                                                                                                                          TISSUE=Venom;
                                                                                                                                                                                                                                                                                                                                     PROTEIN SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                         WADDLINE=87293024; PubMed=3617088; DOI=10.1016/0041-0101(87)90288-1; Yasuhara T., Mantel P., Nakajima T., Piek T.; "Two kinins isolated from an extract of the venom reservoirs of the solitary wasp Megascolia flavifrons.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE-Venom;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROTEIN SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=7437;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neoptera; Endopterygota; Hymenoptera; Scoliidae; Megascolia.
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROTEIN SEQUENCE, MEDLINE=67179312;
                                                                                                                                 -:- SUBCELLULAR LOCATION: Secreted; venom reservoirs.
-:- SIMILARITY: Belongs to the bradykinin family.
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Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
Phyllomedusinae; Phyllomedusa.
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as long as its content in its institute.
                                                                                                                                                            FUNCTION: Both proteins have bradykinin-like, activities (e.g. smooth muscle contraction). SUBCELLULAR LOCATION: Secreted; venom reservo.
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Apocrita; Aculeata;
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Hylidae;
                 restrictions on its
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Q7M4P1;
 PROTEIN SEQUENCE.
MEDLINE=91166748; PubMed=2076202;
Wunderer G., Walter I., Eschenbac
                                                                                              01-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
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PIR; B26744; B26744.

GO; GO:0005615; C:extracellular space;

GO; GO:0045776; P:negative regulation of

GO; GO:0045987; P:positive regulation of
                                                          Mammalia;
                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                    Oncogene 16:3243-3252(1998).
EMBL; U88358; AAC39858.1; -;
                                                                                       Ile-Ser-bradykinin
                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                         TISSUE=Ovarian carcinoma;
MEDLINE=98345147; PubMed=9681822;
Lee H., Maihle N.J.;
"Isolation and characterization of
                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human)
Eukaryota; Metazoa;
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HUMAN
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etazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Euarchontoglires; Primates;
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Pred. No.
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Bradykinin-like peptide.
33867393D771A9C8 CRC64;
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Pred. No.
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VESMA STANDARD; PRT; 12 AA.
VESP VESMA STANDARD; PRT; 12 AA.
Q'MST3;
05-JUL-2004 (Rel. 44, Created)
05-JUL-2004 (Rel. 44, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kindermann G.;
"Ile-Ser-bradykinin is an aberrant permeability factor in various human malignant effusions.";
Biol. Chem. Hoppe-Seyler 371:977-981(1990).
DIR; S13279; S13279.
NON_TER 1 1 1
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                                                                                                                                                                                  MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Meoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Vespoidea; Vespidae; Vespinae; Vespa.

NCBI TaxID=7446;
[1]
                                                                                                                                                                                                                                                                                       This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=77114342; PubMed=1017116;
Kishimura H., Yasuhara T., Yoshida H., Nakajima T.;
Kishimura H., Yasuhara T., Yoshida H., Nakajima T.;
"Vegpakinin-M, a novel bradykinin analogue containing hydroxyproline, in the venom of Vespa mandarinia Smith.";
Chem. Pharm. Bull. 24:2896-2897(1976).
-!- FUNCTION: Bradykinins are a potent but short-lived agent of arteriolar dilation and increased capillary permeability (By
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                               Vasodilator.
                                                                                                                                                                                                                              Bradykinin; Direct protein sequencing; Hydroxylation; Vasoactive;
                                                                                                                                                                                                                                               PIR; A61360; A61360.
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34F513C44C75B9C8 CRC64;
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